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233 13.4 74.4 491 3US- 234 13.4 74.4 530 3 US- 235 13.4 74.4 530 3 US- 236 13.4 74.4 530 3 US- 237 13.4 74.4 601 3 US- 238 13.4 74.4 601 3 US- 239 13.4 74.4 601 3 US- 241 13.4 74.4 601 3 US- 242 13.4 74.4 601 3 US- 243 13.4 74.4 658 3 US-	224 13.8 76.7 421522 3 US-09-949-011 225 13.8 76.7 451924 3 US-09-949-011 226 13.8 76.7 451925 3 US-09-949-011 227 13.8 76.7 451925 3 US-09-949-011 228 13.8 76.7 451726 3 US-09-949-011 229 13.8 76.7 481115 3 US-09-949-011 230 13.8 76.7 1230025 3 US-09-949-011 231 13.8 76.7 1230025 3 US-09-9438-11 231 13.8 76.7 1230230 3 US-09-438-11 232 13.4 74.4 312 3 US-09-513-99	214 13.8 76.7 223471 3 US-09-949-011 215 13.8 76.7 227391 3 US-09-949-011 216 13.8 76.7 227391 3 US-09-949-011 217 13.8 76.7 246240 2 US-08-724-39- 218 13.8 76.7 246240 2 US-08-724-39- 219 13.8 76.7 246240 2 US-08-724-39- 219 13.8 76.7 246240 2 US-08-9249-011 220 13.8 76.7 246243 3 US-09-949-011 221 13.8 76.7 265038 3 US-09-949-011 222 13.8 76.7 300402 3 US-09-949-011 223 13.8 76.7 300402 3 US-09-949-011	177 13.8 76.7 53337 3 US-09-949-011 180 13.8 76.7 55328 3 US-09-949-011 181 13.8 76.7 55328 3 US-09-949-011 181 13.8 76.7 56448 3 US-09-949-011 182 13.8 76.7 66213 3 US-09-949-011 182 13.8 76.7 66213 3 US-09-949-011 183 13.8 76.7 70308 3 US-09-949-011 185 13.8 76.7 70308 3 US-09-949-011 187 13.8 76.7 70308 3 US-09-949-011 187 13.8 76.7 87734 3 US-09-949-011 187 13.8 76.7 19133 3 US-09-949-011 188 13.8 76.7 19133 3 US-09-949-011 191 13.8 76.7 103025 3 US-09-949-011 191 13.8 76.7 103025 3 US-09-949-011 194 13.8 76.7 110025 3 US-09-949-011 195 13.8 76.7 110025 3 US-09-949-011 195 13.8 76.7 112814 3 US-09-949-011 199 13.8 76.7 1134008 3 US-09-949-011 199 13.8 76.7 1134008 3 US-09-949-011 190 13.8 76.7 174318 3 US-09-949-011 13.8 76.7 1205044 3 US-09-949-011 13.8 76.7 205044 3 US-09-949-011 13.8 76.7 2050	171 13.8 76.7 44120 3 US- 172 13.8 76.7 44120 3 US- 173 13.8 76.7 44120 3 US- 174 13.8 76.7 44676 3 US- 174 13.8 76.7 51967 3 US- 175 13.8 76.7 51967 3 US-
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234 13.4 74.4 491 3 US-09-513-9945-2326 235 13.4 74.4 530 3 US-10-012-542-28 235 13.4 74.4 530 3 US-10-012-542-28 236 13.4 74.4 530 3 US-10-115-122-28 237 13.4 74.4 631 3 US-09-949-016-25293 238 13.4 74.4 631 3 US-09-949-016-52531 238 13.4 74.4 631 3 US-09-949-016-12425 240 13.4 74.4 631 3 US-09-949-016-12425 241 13.4 74.4 631 3 US-09-949-016-12425 242 13.4 74.4 631 3 US-09-949-016-151425 243 13.4 74.4 658 3 US-09-949-016-151425 244 13.4 74.4 658 3 US-09-533-559-6647	224 13.8 76.7 422522 3 US-09-949-016-12896 226 13.8 76.7 451924 3 US-09-949-016-12896 227 13.8 76.7 451925 3 US-09-949-016-17305 228 13.8 76.7 451925 3 US-09-949-016-14157 229 13.8 76.7 451915 3 US-09-949-016-11940 229 13.8 76.7 481115 3 US-09-949-016-11940 230 13.8 76.7 1230025 3 US-09-949-016-198-452A-1 231 13.8 76.7 1230203 3 US-09-438-185A-1 232 13.4 74.4 312 3 US-09-513-999C-35050	214 13.8 76.7 223471 3 US-09-949-016-12725 215 13.8 76.7 227390 3 US-09-949-016-12301 216 13.8 76.7 227391 3 US-09-949-016-13365 217 13.8 76.7 246240 2 US-08-724-394A-20 218 13.8 76.7 246240 2 US-08-724-394A-21 219 13.8 76.7 246240 2 US-08-724-394A-22 210 13.8 76.7 246240 2 US-08-724-394A-22 211 13.8 76.7 246240 3 US-09-949-016-13113 220 13.8 76.7 256338 3 US-09-949-016-13632 221 13.8 76.7 300402 3 US-09-949-016-15371 222 13.8 76.7 304533 3 US-09-949-016-15371	176 13.8 76.7 53337 3 US-09-949-016-1294 13.8 76.7 55328 3 US-09-949-016-1294 13.8 76.7 55328 3 US-09-949-016-1294 13.8 76.7 56448 3 US-09-949-016-1244 13.8 76.7 66213 3 US-09-949-016-1264 13.8 76.7 66213 3 US-09-949-016-1264 13.8 76.7 69324 3 US-09-949-016-1264 13.8 76.7 70308 3 US-09-949-016-1264 13.8 76.7 70308 3 US-09-949-016-1264 13.8 76.7 70308 3 US-09-949-016-1264 13.8 76.7 87792 3 US-09-949-016-1264 13.8 76.7 87792 3 US-09-949-016-1264 13.8 76.7 91933 3 US-09-949-016-1264 13.8 76.7 103750 3 US-09-949-016-1264 13.8 76.7 103750 3 US-09-949-016-1264 13.8 76.7 103750 3 US-09-949-016-1264 13.8 76.7 110525 3 US-09-949-016-1264 13.8 76.7 110525 3 US-09-949-016-1264 13.8 76.7 110525 3 US-09-949-016-1265 13.8 76.7 110525 3 US-09-949-016-1265 13.8 76.7 110525 3 US-09-949-016-1265 13.8 76.7 1135010 3 US-09-949-016-1265 13.8 76.7 135010 3 US-09-949-016-1265 13.8 76.7 134008 3 US-09-949-016-1265 13.8 76.7 174170 3 US-09-949-016-1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 1261 13.8 76.7 126	171 13.8 76.7 44120 3 US-09-949-016-14154 172 13.8 76.7 44120 3 US-09-949-016-14155 173 13.8 76.7 44120 3 US-09-949-016-14155 174 13.8 76.7 44676 3 US-09-949-016-17511 175 13.8 76.7 51367 3 US-09-949-016-17582 176 13.8 76.7 51363 1 US-09-949-016-17582

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RESULT 1
US-08-571-758-5
                                                                               Sequence 5, Application US/08571758 Patent No. 5700675
                                           GENERAL INFORMATION:
APPLICANT: Rubin, Gerry M.
APPLICANT: Therrien, Marc
                                  APPLICANT:
                        APPLICANT:
 TITLE OF INVENTION:
                                                                                                                                                                                      Wassarman, David A.
VENTION: A No. 5700675el Protein Kinase Required for Ras
                     Chang, Henry C.
Karim, Felix D.
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US-09-949-016-12544

US-09-949-016-15617
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US-09-949-016-17130
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17433, A
1843, Appl
18544, Appl
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14264,
15999,
12776,
13266,
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46, Appl
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Patent No
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B96-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 343-4341
                                                                                                                                                                                                                                                                                            atent No. 5747275
GENERAL INFORMATION:
                                                                                                                                    APPLICANT: Wassarman, David A.
TITLE OF INVENTION: A No. 5747275el Protein Kinase Required for ITTLE OF INVENTION: Signal Transduction
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
STREET: 268 BUSH STREET, SUITE 3200
CITY: SAN FRANCISCO
STATE: CALIFORNIA
                                                                                                                                                                                                                                             APPLICANT: TAPPLICANT: TAPPLICANT: CAPPLICANT: CAPPLICANT: F
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-BOS #1.0,
                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: US/08/571,758 FILING DATE: CLASSIFICATION: 435
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ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
STREET: 268 BUSH STREET, SUITE 3200
CITY: SAN FRANCISCO
STATE: CALIFORNIA
 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
             FILING DATE:
CLASSIFICATION:
                                  APPLICATION NUMBER:
                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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Chang, Henry C.
Karim, Felix D.
                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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100.0%; Pred. No.
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REGISTRATION NUMBER: 36, REFERENCE/DOCKET NUMBER:

OSMAN,

RICHARD A

36,627

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Sequence 5, Application US/08909983 Patent No. 5747288
                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B96-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 343-4341
TELEPHONE: (415) 343-4342
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/909,983
FILING DATE: 12-JUN-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/571,758
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TELEPHONE: (415) 343-4341
TELEFAX: (415) 343-4342
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Wassarman, David A.
TITLE OF INVENTION: A No. 5747288el Prof
TITLE OF INVENTION: Signal Transduction
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: CDNA
MOLECULE TYPE: cDNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP STREET: 268 BUSH STREET, SUITE 3200 CITY: SAN FRANCISCO
                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                    STRANDEDNESS:
                   TOPOLOGY:
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                                                                                ENGTH:
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                                                                              4094 base pairs
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Karim, Felix D
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Pred. No. 5
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Query Match

100.0%;

Score 18;

DB 2;

Length 4094;

US-09-949-016-12212/c

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                                                                                                                            ; ORGANISM: Human US-09-949-016-13682
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                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 13682
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: VENTER,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13682, Application US/09949016 Patent No. 6812339
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Patent No. 6639063
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                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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CURRENT FILING DATE: 2000-07-21
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                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
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                                                                                                                                                            TYPE: DNA
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                                                                                                                                                                           ENGTH: 29889
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                                                                              Local
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Local Similarity 94.1%;
29541 GGÁCCCAAGAGCCAAAG 29557
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                              2 GGACCCTAGAGGCAAAG 18
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                                                                              Similarity
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Pred. No. 1
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Pred. No. 1
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; LOCATION: (1)...(97196)
; OTHER INFORMATION: n = A,T,C
US-09-949-016-16971
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; LICATION: (1)...(97195)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-12212
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SEQ ID NO 16971
LENGIH: 97196
                                                                                                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR PILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
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CURRENT FILING DATE: 2000-04-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: CL001307
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
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CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                            ORGANISM: Human FEATURE:
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ORGANISM: Human
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                                                                                                    Similarity
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Similarity 94.1%;
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GGACCCTAGAGGCAAGG 77954
                                      GGACCCTAGAGGCAAAG 18
                                                                              Conservative
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94.1%;
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Pred. No. 1.7e+02;
                                                                                                  Score 15.4; DB 3;
Pred. No. 1.7e+02;
                                                                            Mismatches
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                                                                                                                     Length 97196;
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                                                                                                                                               ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-513-999C-28900
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; ORGANISM: Homo sapiens
US-10-131-827-7589
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US-10-131-827-7589
                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 28900
                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins Patent No. 6783961
FILE REFERENCE: 59.US2.REG
                                                                          Matches
                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 28900, Application US/09513999C Patent No. 6783961
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Best Local
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                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y.
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CURRENT FILING DATE: 2002-09-06
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/296,764 PRIOR FILING DATE: 2001-06-08
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APPLICANT: Fry, Kirk
APPLICANT: Woodward, Robe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 9090
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING AUTOIMMITTE OF INVENTION: CHRONIC INFLAMMATORY DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 50
                                                                                                                                                                                                      ENGTH: 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
148 GGACCCTAGAGGCAA 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                        15;
                                   2 GGACCCTAGAGGCAA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGACCCTAGAGGCAA 32
                                                                                                                                                                                                                                                                                                                                                                                                              Giordano,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fry, Kirk
Woodward, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ly, Ngoc
                                                                        Conservative
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                                                                                            100.0%;
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                                                                        0;
                                                                                        Score 15;
Pred. No.
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                                                                          Mismatches
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                                                                                            1.5e+02
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RESULT 10 US-09-157-910-6

GENERAL INFORMATION:

Sequence 6, Application US/09157910A Patent No. 6197744

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RESULT 12
US-09-023-655-508
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                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PE
SEQ ID NO 159
LENGTH: 868
                                               Sequence 508, Application US/09023655 Patent No. 6607879
                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6426186
GENERAL INFORMATION:
APPLICANT: Jones,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 159,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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LENGTH: 783
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/484,970B
CURRENT FILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/157,910A
CURRENT FILING DATE: 1998-09-21
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Microsoft Word 97 / MS-DOS Ver. 7.10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-484-970B-159
                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Tumor Necrosis Factor Inhibitory Protein TIP-B1 and Method TITLE OF INVENTION: of Using Same FILE REFERENCE: RPP:151-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TILE REFERENCE:
                                                                                                                                                                                                                                                                                               NAME/KEY: unsure LOCATION: 23, 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE: OTHER INFORMATION: This sequence encodes the TIP-B1 peptides of SEQ. OTHER INFORMATION: SEQ. ID. #4 and SEQ. ID. #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: HOMO SAPIENS
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PPLICANT: Mihich, Enrico
                                                                                                                                                                                                                                                                                   OTHER INFORMATION: a,
                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. 6426186 234571.8
                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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APPLICANT: Cocks, Benjamin G. APPLICANT: Susan G. Stuart
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les 15; Conserv
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100.0%; Pred. No. 1.8e+02;
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RESULT 13
US-09-949-016-23074/c
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                                                                 CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                  Sequence 23074, Application US/09949016 Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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SEQ ID NO 23074
LENGTH: 601
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INFORMATION FOR SEQ ID NO: 508:
SEQUENCE CHARACTERISTICS:
                               SOFTWARE:
                                                  NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                              SMERAL INFORMATION:

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                   SNERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA: US
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for 1
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CLONE: 201392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
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Similarity 100.0%; F
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                                 FastSEQ for Windows Version 4.0
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3174 PORTER DRIVE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HEREWITH
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; Pred. No. 2e+
0; Mismatches
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; ORGANISM: Human
US-09-949-016-169781
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                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Appl Patent No. 57006
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Best Local Similarity
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: CL001307
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                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                TITLE OF INVENTION: A No. 5700675el Protein Kinase Required for Ras
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APPLICATION NUMBER:
                                                                                                                                                                                   STREET:
                                                                                                                                COUNTRY:
                                                                                                                                                                                                   DDRESSEE:
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                                                                                                                   94104
                                                                                                                                                 SSEE: SCIENCE & TECHNOLOGY LAW GROUP
1: 268 BUSH STREET, SUITE 3200
SAN FRANCISCO
: CALIFORNIA
                 APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                       Application US/08571758
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                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                   Wassarman,
                                                                                                                                                                                                                                                                                                 Chang, Henry C.
Karim, Felix D.
                                                                                                                                                                                                                                                                                                                                      Therrien, Marc
                                                                                                                                                                                                                                                                                                                                                      Rubin, Gerry M.
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US/08/571,758
                                                                                                                                                                                                                                                                                   David A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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Pred. No. 2.2e+02;
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Pred. No. 2.2e+02
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US-08-909-984A-7
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                                                                                              TELEPHONE: (415) 343-43
TELEFAX: (415) 343-4342
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                             FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REGISTRATION NUMBER: 36,627
                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/909,984A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                   TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Wassarman, David A.
TITLE OF INVENTION: A NO. 5747275el Prot
TITLE OF INVENTION: Signal Transduction
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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NAME: OSMAN, RICHARD A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 343-4341
                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
STREET: 268 BUSH STREET, SUITE 3200
CITY: SAN FRANCISCO
                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
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CLASSIFICATION:
                 COPOLOGY:
                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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                                                                 ENGTH:
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                                          nucleic acid
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TYPE: CDNA
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Karim, Felix D.
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              linear
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                                 double
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88.9%;
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Pred. No. 2.
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Query Match

82.2%;

Score 14.8;

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2

Length 2846;

Matches

Best Local Similarity

ilarity 88.9%; Conservative

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Pred. No.

ed. No. 2.5e+02; Mismatches 2;

Indels

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Gaps

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RESULT 18
US-09-560-385A-23
; Sequence 23, Application US/09560385A
; Patent No. 6703363
; Patent No. 6703363
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US-08-909-983-7
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                                                                                                                                                                                                                                     Matches
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Best Local Similarity
                 APPLICANT: Boutand, CITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (415) 343-4342 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
   ILE REFERENCE:
                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: B9
TELECOMMUNICATION INFORMATION
TELEPHONE: (415) 343-4341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/909,983
FILING DATE: 12-UUN-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: À NO. 5747288el Proi
TITLE OF INVENTION: Signal Transduction
UMBER OF SEQUENCES: 12
ORRESPONDENCE ADDRESS:
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VENTION: A No. 5747288el Protein Kinase Required for Ras
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Karim, Felix D.
ON: Recombinant Laminin 99-274-C
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                                    Ariel
                                                                                                                                                                                                                                                   82.2%;
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Pred. No. 2
                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                      DB
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                                                                                                                                                                                                                                   Sequence 21, Application US/09560385A Patent No. 6703363
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Best Local Similarity
                                                                                           SOFTWARE:
SEQ ID NO 21
                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             Query Match 82.3
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19, Application US/09560385A
Patent No. 6703363
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                                                                                                                                         FILE REFERENCE: 99-274-C
CURRENT APPLICATION NUMBER: US/09/560,385A
CURRENT FILING DATE: 2000-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/560,385A
CURRENT FILING DATE: 2000-04-28
CUMBER OF SEQ ID NOS: 36
                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                  APPLICANT: Boutand, Ariel
TITLE OF INVENTION: Recombinant Laminin 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE:
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CURRENT FILING DATE: 2000-04-28
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TITLE OF INVENTION: Recom
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                                                              LENGTH: 3585
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (7)..(3507)
NAME/KEY: misc_feature
LOCATION: (13)..(36)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 3469
TYPE: DNA
FEATURE:
NAME/KEY: CDS
LOCATION: (60)..(3581)
                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                  1858 CGGATCCTAGATGCAAAG 1875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3511
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                                                                                                            PatentIn Ver. 2.0
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Pred. No. 2
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Pred. No. 2.6e+02;
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RESULT 22
US-09-560-385A-15
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US-09-560-385A-17
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; LOCATION: (60)..(116)
US-09-560-385A-21
                                                                              JS-09-560-385A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-560-385A-17
     Query Match
Best Local Similarity 85.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: P
SEQ ID NO 17
LENGTH: 362
TYPE: DNA
                                                                                                                                                                       SOFTWARE: 1
SEQ ID NO 15
                                                                                                                                                                                                                                                                                            Sequence 15, Application US/09560385A
                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                        TILE REFERENCE: 99-274-C
CURRENT APPLICATION NUMBER: US/09/560,385A
CURRENT FILING DATE: 2000-04-29
NUMBER OF 7-7-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (60)..(3617)
NAME/KEY: SIG DEPLIGE
LOCATION: (60)..(110)
NAME/KEY: misc feature
LOCATION: (123)..(146)
                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/560,385A
CURRENT FILING DATE: 2000-04-28
NUMBER OF SEQ ID NOS: 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Boutand, Ariel
TITLE OF INVENTION: Recom
                                                                                                                     LENGTH: 3759
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                        NAME/KEY: CDS
LOCATION: (1)..(3459)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                    PPLICANT: Boutand, Ariel
ITLE OF INVENTION: Recombinant Laminin 5
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NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                           1968 CGGATCCTÁGATGCAAAG 1985
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1 CGGACCCTAGAGGCAAAG
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                                     Score 14.8;
Pred. No. 2
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Pred. No. 2.
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                                    2.6e+02;
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                                                  DB 3;
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                                                                                                                                                              CURRENT APPLICATION NUMBER: 27-007-1993

CLASSIFICATION: 424

ATTORNEY / ACTIVATION: ALCANIA CARREL 27-007-1993

CLASSIFICATION: 424

ATTORNEY / ACTIVATION: 424
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        ATTORNEY/AGENT INFORMATION:
NAME: Myers, Paul L.
REGISTRATION NUMBER: 35,965
REFERENCE/DOCKET NUMBER: (MGH-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 3931 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1,
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APPLICANT: Burgeson, Robert E.
APPLICANT: Wagman, David W.
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CURRENT FILING DATE: 2000-C
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (121)..(3630)
NAME/KEY: sig_peptide
LOCATION: (121)..(171)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
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3931 base pairs
ucleic acid
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88.9%;
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Pred. No. 2
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US-08-735-893-1
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Query Match 82. Best Local Similarity 88. Matches 16; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                  TELEFAX: (617) 227-5941
.NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 3931 base pairs
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APPLICATION NUMBER: US 08/144,121
FILING DATE: 27-OCT-1993
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                            MOLECULE TYPE:
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                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: .(617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY:
LOCATION:
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                                                                                                                 NAME/KEY:
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                                                                                                                                                                                                                                                                                                       NAME: Myers, Paul L.
REGISTRATION NUMBER: 35,965
REFERENCE/DOCKET NUMBER: (M
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/0 FILING DATE: 18-OCT-1996 CLASSIFICATION: 43'5
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TOPOLOGY: lir
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milarity 88.9%;
Conservative
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            82.2%;
88.9%;
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Score 14.8; DB 2;
Pred. No. 2.6e+02;
0; Mismatches 2;
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Pred. No. 2
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                                                                                                       ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-876-176-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: mat peptide; LOCATION: (173)..(3616)
US-10-841-139-1
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                                    Matches
                                                  Query Match
Best Local Similarity
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Best Local
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APPLICANT: Winter, Jill A.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING
TITLE OF INVENTION: NEOPLASTIC DISEASE USING INHIBITORS OF LAMININSBETA3
FILE REFERENCE: PP-01702.002/200130.523
CURRENT APPLICATION NUMBER: US/09/876,176
CURRENT FILING DATE: 2001-06-06
NUMBER OF SEO ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (122)..(3616)
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CURNENT FILING DATE: 2004-05-07
FRIOR APPLICATION NUMBER: US/10/443,349
PRIOR FILING DATE: 2003-05-22
PRIOR APPLICATION NUMBER: US/09/161,872
PRIOR FILING DATE: 1998-09-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Burgeson, Robert E. APPLICANT: Wagman, David W.
                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0 EQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver.
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APPLICANT: Jefferson, Anne B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 10287/021003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wagman, David W.
FITLE OF INVENTION: Blk CHAIN OF LAMININ AND METHODS OF USE
                                                                                                                                                            ENGTH:
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   CGGACCCTAGAGGCAAAG 18
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Den-Otter, Douglas R.
                                      Conservative
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88.9%;
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Pred. No. 2.6e
0; Mismatches
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Pred. No. 2
                                      Mismatches
                                                        2.6e+02;
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                                                                        Length 3971;
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US-07-841-646-3
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PERL Program
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Turner, Christopher M.
TITLE OF INVENTION: PROSTATE CANCER MARKERS
FILE REFERENCE: PA-0.036 US
CURRENT APPLICATION NUMBER: US/09/919,172
CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/222,469
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 102
                                          APPLICATION NUMBER: US/07/841,64
FILING DATE: 19920221
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 810,560
FILING DATE: 20-DEC-1991
PRIOR APPLICATION DATA:
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APPLICANT: Turner, Ch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. 6673545 3615080CB1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: OSTEOGENIC DEVICES NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS: TESTA, HURWITZ & THIBEAULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH: 4108
                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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              APPLICATION NUMBER: FILING DATE: 28-JAN
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APPLICATION DATA:
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RUEGER, DAVID C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPPERMANN, HERMANN
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            28-JAN-1992
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                            US 827,052
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Pred. No. 2.6e+02;
0; Mismatches 2
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17410 base pai
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LOCATION: 10413.10414
OTHER INFORMATION: /labe
OTHER INFORMATION: /not
OTHER INFORMATION: BE M.
                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                          NAME/KEY: exon
LOCATION: 3192..3730
OTHER INFORMATION: /Lo
OTHER INFORMATION: /no
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PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:

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APPLICATION NUMBER: US 616,374
FILING DATE: 21-NOV-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
PRIOR APPLICATION UTMBER: US 599,543
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PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
LOCATION: 10960..10961
OTHER INFORMATION: /label= GAP-2
                                                                      LOCATION: 10696..1
OTHER INFORMATION:
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                                                                                                                                           OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
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LECOMMUNICATION INFORMATION:
TELEPHONE: 617/248-7000
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                     NAME/KEY: misc_feature
                                                                                                         NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                     ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 17410 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                                                    10696..10891
RMATION: /label= EXON-2
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23-FEB-1989
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22-FEB-1990
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20-AUG-1990
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07-SEP-1990
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                                                                                                                                        /note= "APPROXIMATELY BASES ARE ESTIMATED BE MISSING BETWEEN POSITIONS 10413 AND 10414 IN THIS SEQUENCE."
                                                                                                                                                                                                                                                               /label= EXON-1
/note= "START CODON BEGINS AT POSITION 3313"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US 179,460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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URRENT APPLICATION DATA
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LOCATION: 11721..11722

OTHER INFORMATION: /label= GAP-4

OTHER INFORMATION: /note= "APPROXIMATELY BASES ARE ESTIMATED TO OTHER INFORMATION: BE MISSING BETWEEN POSTTTONS 11721 AND 11722 IN OTHER INFORMATION: THIS SEQUENCE."
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OTHER INFORMATION: /label= EXON-7
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LOCATION: 11059..11211
OTHER INFORMATION: /la
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                                                                                                                                                               STREET:
                                                                                                                                                                                                                                    TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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LOCATION: 11721..11722
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LOCATION: 11420..11617
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                                                                                                              COUNTRY:
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LOCATION: 11351..11352
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                                                                                                                                MASSACHUSETTS
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OZKAYNAK, ENGIN
KUBERASAMPATH, TH
RUEGER, DAVID C.
                                                                                                                                                               E: TESTA, HURWITZ & THIBEAULT
53 STATE STREET:
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RMATION: /label= EXON-6
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/note= "APPROXIMATELY BASES ARE ESTIMATED TO BE MISSING BETWEEN POSITIONS 11351 AND 11352 IN THIS SEQUENCE."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "APPROXIMATELY BASES ARE ESTIMATED BE MISSING BETWEEN POSTTION 10960 AND 10961 IN THIS SEQUENCE."
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                                                                                                                                                                                                                                                                                                                    HERMANN
                                                                                                                                                                                                                                                                                   THANGAVEL
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Pred. No. 3e+0
O; Mismatches
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3e+02;
               #1.25
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TELEFAX: 617/248-7100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 170
PILING DATE: 00
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                                                                                                                                                                  MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/248-7000
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ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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NAME/KEY: exon

LOCATION: 15044..15160

OTHER INFORMATION: /label= EXON-6

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NAME/KEY: exon

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/note= "APPROXIMATELY BASES ARE ESTIMATED TO BE MISSING BETWEEN POSITIONS 11721 AND 11722 IN THIS SEQUENCE."
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Aai21300
Aba68190
Aba66383
Aba66383
Aba66389
Aai447027
Aai46576
Aai446940
Aba48940
Aba55198
Aba55198
Aba33444
                                                                        Aak15248
Aak14798
Aak16565
Abs40102
Abs41936
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Aai88464
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Abs16285
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Abs41841
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Aai35382
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                                    9 Lymphona
3 Human foe
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Human foe
1 Probe #24
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Jobe #12
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                         The invention comprises antisense oligonucleotides which are designed to inhibit the expression of the kinase suppressor of Ras (KSR) gene. The antisense oligonucleotides of the invention are useful for treating or inhibiting the progression of cancer, such as: pancreatic cancer, lung cancer, skin cancer, urinary tract cancer, bladder cancer, liver cancer thyroid cancer, colon cancer, intestinal cancer, breast cancer, ovarian cancer, stomach cancer, head and neck cancer, because cancer, cancer, lung prostate cancer, leukaemia, lymphoma, and neuroblastoma. The present DNA sequence represents a human KSR-specific oligonucleotide that was used in an example of the invention.
                                                                                                                                                                                                                                                                                         Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense oligonucleotides complementary to a region of kinase suppressor of Ras (KSR) RNA which inhibit KSR expression, useful for treating or inhibiting progression of cancer, e.g. pancreatic, lung, skin, or bladder cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; SEQ ID NO 15; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-053335/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kolesnick RN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-2002; 2002US-0384228P 03-APR-2003; 2003US-0460023P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense oligonucleotide; kinase suppressor of Ras; KSR; cancer; pancreatic cancer; lung cancer; skin cancer; urinary tract cancer; bladder cancer; liver cancer; thyroid cancer; colon cancer; intestinal cancer; breast cancer; ovarian cancer; stomach cancer; intestinal cancer; breast cancer; ovarian cancer; stomach cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    intestinal cancer; breast cancer; ovarian cancer; stomach cancer;
head cancer; neck cancer; oesophageal cancer; prostate cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human kinase suppressor of Ras (KSR)-specific oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lymphoma; neuroblastoma; ss; human.
                                                                                                                                                                                                                18;
                                                                                                                                                                                                                                  Similarity
                                    standard;
                                                                                                                                                           CGGACCCTAGAGGCAAAG 18
                                                                                                                                     CGGACCCTAGAGGCAAAG
                                                                                                                                                                                                                                                                                         BP; 6 A; 5 C; 6 G;
                                                                                                                                                                                                                Conservative
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                                      DNA;
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Pred. No. 21;
0; Mismatches
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RESULT 1
ADG75398
ID ADG7
XX
AC ADG7
XX ADG75398; ADG75398 standard; 18 ₽₽

ALIGNMENTS

12; <u>.</u>

Length 18; Indels

in

Gaps

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RESULT 2
ADG75391;
XX ADG75391;
XC ADG75391;
XC ADG75391;
XX T1-MAR-200
XX T1-MAR-200
XX T1-MAR-200
XX DE Human kina
XX Intisense
XX Dancreati
XX Dancreat pancreatic cancer; lung cancer; skin cancer; urinary tract bladder cancer; liver cancer; product cancer; colon cancer; cintestinal cancer; breast cancer; ovarian cancer; stomach chead cancer; neck cancer; ossophageal cancer; prostate canc Human kinase suppressor of Ras (KSR) antisense oligonucleotide #3 11-MAR-2004 antisense oligonucleotide; kinase suppressor of Ras; lymphoma; neuroblastoma; ss; human. (first entry) cancer; urinary tract cancer; KSR; cancer; cancer; cancer; leukaemia;

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Best Local
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                                                                                                           gastrointestinal disease; respiratory disease; genitourinary disease; endocrine disease; bladder tumor; colon tumor; intestine tumor; cheen tumor; colon tumor; intestine tumor; head & neck tumor; leukemia; hematological disease; lymphoma; immune disorder; esophagus tumor; breast tumor; stomach tumor; nervous system tumor; neurological disease, ovary tumor; synecology and obstetrics; ss; angiogenesis disorder; antiangiogenic; cardiovascular disease; antisense oligonucleotide; antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides complementary to a region of kinase suppressor of Ras (KSR) RNA which inhibit KSR expression, useful for treating or inhibiting progression of cancer, e.g. pancreatic, lung, skin, or bladder cancer.
                                                                                                                                                                                                                           Kinase suppressor of Ras; KSR; cytostatic; neoplasm; pancreas tumor; liver tumor; skin tumor; thyroid tumor; lung tumor; urinary tract tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 8; SEQ ID NO 8; 120pp; English
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03-APR-2003;
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                                                                                                                                                                                                                                                                             Kinase suppressor of Ras (KSR) antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                             25-AUG-2005
                                                                                                                                                                                                                                                                                                                                             AEA46548;
                                                                                                                                                                                                                                                                                                                                                                          AEA46548 standard; DNA; 18
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                                                                                             prostate tumor;
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2003US-0460023P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003WO-US016961
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                                                                                             andrology;
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                                                                                             cancer; hyperproliferation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 18; DB
Pred. No. 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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New oligonucleotide inhibiting kinase suppressor of Ras (KSR) expression, useful for inhibiting expression of mammalian KSR and treating
                                                                                                                                                                                                                                                           WPI; 2005-445165/45
                                                                                                                                                                                                                                                                                    03-DEC-2003; 2003US-00727358
                                                                                                                                                                                                                                                                                            03-DEC-2004; 2004WO-US040506
                                                                                                                                                                                                                                           hyperproliferative conditions such as lung cancer, skin cancer,
                                                                                                                                                                                                                                                                    Kolesncik RN,
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Example 2; SEQ ID NO 15; 186pp; English

cc sequence substantially complementary to nucleotides (14) comprising a sequence of mouse KSR or nucleotides 97-216 of human KSR or its portion; an antisense oligonucleotide (V) comprising a sequence chosen from ARA4539, ARA6540 and ARA6541, and; a recombinant DNA molecule (VI) comprising a nucleic acid sequence with encodes on transcription and antisense NAA complementary to memmalian KSR RNA or its portion; a cell (VII) transfected with (VII); an expression vector (VI) capable of expressing a nucleic acid which is substantially complementary to the coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide/nucleic acid inhibits the expression vector (VI) capable of KSR RNA, or its portion/fragment, where the coding sequence of KSR RNA, or its portion/fragment where the coding memmalian KSR and which is substantially complementary to the CSR RNA, or its portion/fragment, where the coding sequence of KSR, and, or its portion/fragment, where the coding sequence of KSR, and, or its portion/fragment, where the coding sequence of CSR RNA, or its portion/fragment, where the cligonucleotide which is the coding sequence of compression of KSR, a pharmaceutical composition (PCI) comprising (I) and a carrier or diluent, a composition (CI) comprising one or more chemotherapeutic or radiotherapeutic agent and an oligonucleotide which is targeted to a mRNA coding memmalian KSR and which inhibits the expression; a composition or radiotherapeutic or substantially complementary to the coding sequence of KSR RNA, or its portion/fragment, where the nucleic acid which is composition of KSR; treating or preventing (MZ) a portion of the mRNA encoding KSR; treating or preventing (MZ) a preventing (MZ) a preventing (MZ) a preventing to the mammal acomposition of mammalian KSR protein, inhibits the expression of Ras i The invention relates to an oligonucleotide (I) which is substantially complementary to a region of kinase suppressor of Ras (KSR) RNA, where the oligonucleotide inhibits the expression of KSR. Also included are the following: an oligonucleotide (II) which is substantially complementary to a translation initiation site, 5' untranslated region, coding region or 3' untranslated region of mRNA encoding mammalian KSR; an antisense oligonuclectide (III) comprising a sequence substantially complementary to the CA1 region of KSR; an antisense oligonuclectide (IV) comprising the expression of mammalian KSR protein or a vector expressing mammalian KSR; identifying compounds or agents which inhibit the expression of KSR by incubating a cell expressing KSR in the presence and absence of a angiogenesis in a mammal comprising administering to a mammal a therapeutically effective amount of a compound or agent which activates

WO2005056756-A2

Homo sapiens

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RESULT 4

REA46538

ID ABA4

XX AEA4

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kinase suppressor of Ras; KSR; cytostatic; neoplasm; pancreas tumor; liver tumor; skin tumor; thyroid tumor; lung tumor; urinary tract tumo gastrointestinal disease; respiratory disease; genitourinary disease; endocrine disease; bladder tumor; colon tumor; intestine tumor; head & neck tumor; leukemia; hematological disease; lymphoma; immune disorder; esophagus tumor; breast tumor; stomach tumor; prevous system tumor; neurological disease; unor; stomach tumor; synecology and obstetrics; ss; angiogenesis disorder; antisense therapy; prostate tumor; andrology; cancer; hyperproliferation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kinase suppressor of Ras (KSR) antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         candidate compound or agent, and detecting or measuring the expression of KSR in the presence and absence of a candidate compound or agent, where a decrease in the expression of KSR in the presence of the candidate compound or agent versus in the absence of the candidate compound or agent indicates that the compound or agent inhibits the expression of KSR agent indicates that the compound or agent inhibits the expression of KSR and a ribozyme that cleaves KSR mRNA. (I) Is useful for inhibiting the expression of mammalian KSR comprising contacting cells which express KSR with (I), where expression of mammalian KSR is inhibited. (I) Is useful
                                                                                                                                                                                                                                                                                                                                                                                                                            03-DEC-2004; 2004WO-US040506
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                                                                                                                     WPI; 2005-445165/45
                                                                                                                                                                                             Kolesncik RN,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2005056756-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-AUG-2005
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Pred. No. 21;
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New oligonucleotide inhibiting kinase suppressor of Ras (KSR) expression, useful for inhibiting expression of mammalian KSR and treating

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> ovarian cancer. hyperproliferative conditions such as lung cancer, skin cancer, and

7; SEQ ID NO 5; 186pp; English

C sequence of mouse KSR cor huclectudes 9/-216 of numan KSR cor its portion; an artisense cligonolectide (VI) comprising a sequence chosen from comprising a nucleic acid sequence which encodes on transcription and relations of the motivation of the comprising a nucleic acid sequence which encodes on transcription and relations of the comprising and comprising of the coding sequence of KSR kNA could be presented to the coding sequence of KSR kNA which is expression in the coding sequence of KSR kNA which is expression of KSR; and composition (VI) comprising of the coding sequence of KSR kNA which is purticular transcription of KSR; and composition (VI) comprising (I) and a carrier or diluent, a composition (CI) comprising on an injunct code inhibite the expression of KSR; and coding mammalian KSR and which inhibits KSR expression; a composition (CI) comprising on a coding sequence of KSR kNA, or its portion/fragment, where the coding sequence of KSR kNA, or its composition (CI) comprising (I) and a carrier or diluent, where the coding sequence of KSR kNA, or its conclusion weeter is capable of expression inhibite the expression; a composition (CI) comprising (I) and a carrier or diluent, where the coding sequence of KSR kNA, or its conclusion weeter is capable of expression inhibite the expression of KSR; and which inhibits KSR expression; a composition (CI) comprising uncleic acid which is complementary to a portion of the expression of kRS; inhibiting (NI) the expression of mammalian KSR comprising contacting contacting contacting contacting an expression of the coding sequence of KSR; inhibiting the expression of the mammalian KSR comprising dominant trailing to the mammal comprising dominant trailing contacting to the mammal comprising dominant trailing contacting to the mammal comprising dominant trailing contacting to the mammal comprising that the expression of mammalian KSR protein; trailing to the mammalian KSR protein or a weater expression of the mammalian KSR protein; stimulating creditions of the ex to the CA1 region of KSR; an antisense oligonucleotide (IV) comprising a sequence substantially complementary to nucleotides 124-243 of the coding sequence of mouse KSR or nucleotides 97-216 of human KSR or its portion; to a translation initiation site, 5' untranslated region, coding region of 3' untranslated region of mRNA encoding mammalian KSR; an antisense oligonucleotide (III) comprising a sequence substantially complementary to the CA1 region of KSR; an antisense oligonucleotide (IV) comprising The invention relates to an oligonucleotide (I) which is substantially complementary to a region of kinase suppressor of Ras (KSR) RNA, where the oligonucleotide inhibits the expression of KSR. Also included are titollowing: an oligonucleotide (II) which is substantially complementary cancer in a mammal. The the

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RESULT 5
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ID AEA465
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bladder cancer, liver
cancer, leukemia, lymp
                                                                                                                                                                                                                                                                                                                                                                                                                       New oligonucleotide inhibiting kinase suppressor of Ras (KSR) useful for inhibiting expression of mammalian KSR and treating hyperproliferative conditions such as lung cancer, skin cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SLOAN
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                                                                                                                                                                                                                                                                                                                                                                                                                          cancer, skin cancer, and
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The invention relates to an oligonucleotide (I) which is substantially complementary to a region of kinase suppressor of Ras (KSR) RNA, where the oligonucleotide inhibits the expression of KSR. Also included are the following: an oligonucleotide (II) which is substantially complementary

Claim 8;

SEQ ID NO 28; 186pp; English.

to a translation initiation site, 5' untranslated region, coding region or 3' untranslated region of mRNA encoding mammalian KSR; an antisense

oligonucleotide (III) comprising a sequence substantially complementary to the CA1 region of KSR; an antiense oligonucleotide (IV) comprising a sequence substantially complementary to nucleotides 124-24 of the coding sequence of mouse KSR or nucleotides 97-216 of human KSR or its portion;

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Matches Query Match Best Local &

18;

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Mismatches

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18

Similarity

100.0%;

Score 18; Pred. No.

DB 14;

Length

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ce cells which express Make with a nucleic acid which is complementary to a computencial (M2) and control of the mRNA encoding KSR, treating or preventing (M2) a hyperproliferative condition associated with the expression of gf-Ras or cell heightened expression of Ras in a mammal comprising administering to the associated with the expression of gf-Ras or protein; treating or preventing (M3) a hyperproliferative condition associated with the expression of gf-Ras or heightened expression of Ras in a mammal comprising to the mammal therapeutically effective amount of a nucleic acid which is complementary to a portion of the mRNA encoding KSR, treating or inhibitistering to a mammal a compound or agent which inhibits the expression of cancer in a mammal comprising calministering to a mammal a compound or agent which inhibits the expression of mammalian KSR protein; inhibiting or reducing to the expression of mammalian KSR protein; inhibiting or reducing to agent that inhibits the expression of mammalian KSR protein; stimulating or reducing trace expression of mammal comprising administering to a mammal a compound or agent which national activates the expression of mammal activates of a candidate compound or agent which inhibit the expression of KSR, in the presence and absence of a candidate compound or agent which activates agent inhibiting a cell expression of KSR in the presence of the candidate compound or agent which activates agent which activates agent which inhibit the expression of KSR, in the presence of the candidate compound or agent which activates and a ribozyme that cleaves KSR mRNA. (I) Is useful for inhibiting the expression of faminialian KSR comprising or adiation in tumor cells in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC Comprising a nucleic acid sequence which encodes on transcription and applicated (VI) comprising a nucleic acid sequence which encodes on transcription and antisense RNA complementary to mammalian KSR RNA or its portion; a cell CC (VII) transfected with (VI); an expression vector (VI) capable of coding sequence of KSR RNA, or its portion/fragment, where the coligonucleotide/muclest acid which is substantially complementary to the coligonucleotide/muclest acid inhibits the expression of KSR; an composition of the coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide which is substantially complementary to the CA1 region of the coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide which is substantially complementary to the CA1 region of the coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide inhibits the expression of KSR; a pharmaceutical composition (PC1) comprising (I) and a carrier or diluent; a composition (C1) comprising (I) and a carrier or diluent; a composition (C2) comprising one or more chemotherapeutic or composition (C3) comprising an expression vector and a carrier or diluent, where the expression vector and a carrier or diluent, where the coding sequence of KSR RNA, or its portion/fragment, where the nucleic acid inhibits the expression of KSR; inhibiting (M1) the expression of mammalian KSR comprising contacting cells which express KSR with a nucleic acid which is complementary to a nortice of which is complementary to the coding sequence of KSR RNA, or its portion/fragment, where the nucleic acid inhibits the expression of KSR; inhibiting (M1) the expression of mammalian KSR comprising contacting cells which is expression and a carrier or diluent.
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                                                                                                   ovarian cancer, stomach cancer, The present sequence represents
18
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C; 5
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T; 0 U; 0 Other;
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The invention relates to an oligonucleotide (I) which is substantially CC complementary to a region of kinase suppressor of Ras (KSR) RNA, where the oligonucleotide inhibits the expression of KSR. Also included are the CC following: an oligonucleotide (II) which is substantially complementary to a translation initiation site, 5 untranslated region, coding region (CC or 3' untranslated region of mRNA encoding mammalian KSR; an antisense CC oligonucleotide (III) comprising a sequence substantially complementary to the CA1 region of KSR; an antisense oligonucleotide (IV) complementary CC to the CA1 region of KSR; an antisense oligonucleotide (IV) comprising a sequence substantially comprising a sequence of mouse KSR or nucleotides 97-216 of human KSR or its portion; CC sequence of mouse KSR or nucleotides 97-216 of human KSR or its portion; CC an antisense oligonucleotide (V) comprising a sequence chosen from CC AEA46539, AEA46540 and AEA46541, and; a recombinant DNA molecule (VI) CC comprising a nucleic acid sequence which encodes on transcription and CC (VII) transfected with (VI); an expression vector (VI) capable of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotide inhibiting kinase suppressor of Ras (KSR) expression, useful for inhibiting expression of mammalian KSR and treating hyperproliferative conditions such as lung cancer, skin cancer, and
                  expressing a nucleic acid which is substantially complementary to coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide/nucleic acid inhibits the expression of KSR; an
                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             liver tumor; skin tumor; thyroid tumor; lung tumor; urinary tract tum gastrointestinal disease; respiratory disease; genitourinary disease; endocrine disease; bladder tumor; colon tumor; intestine tumor; head & neck tumor; leukemia; hematological disease; lymphoma; immune disorder; esophagus tumor; breast tumor; stomach tumor; nervous system tumor; neurological disease; ovary tumor; gynecology and obstetrics; ss; angiogenesis disorder; antiangiogenic; gynecology and obstetrics; ss; angiogenesis disorder; antiangiogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kolesncik RN,
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DB; AEA46535.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KETTERING INST CANCER RES
                                                                                                                                                                                                                                                                                                                                                                                                                                      NO'1; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /product= "Kinase suppressor of Ras (KSR) CA1 domain"
/note= "No start or stop codon shown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'partial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
capable of expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              עם
an oligonucleotide which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO:1.
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Conhibiting (MI) the expression of mammalian KSR comprising contacting colls which express KSR with a nucleic acid inhibits the expression of KSR; complementary to a portion of the mRNA encoding KSR; treating or preventing (M2) a hyperproliferative condition associated with the expression of mammalian KSR complementary to a portion of the mRNA encoding KSR; treating or preventing (M2) a hyperproliferative condition associated with the expression of mammalian KSR protein; treating or preventing (M3) a hyperproliferative condition associated with the expression of mammalian KSR complementary to a portion of the mRNA encoding KSR; treating or in a mammal comprising expression of gf-Ras or heightened expression of Ras in a mammal comprising to the complementary to a portion of the mRNA encoding KSR; treating or inhibits the expression of complementary to a portion of the mRNA encoding KSR; treating or inhibits the expression of mammalian KSR protein; stimulating a compound or agent which compound or agent which compound or agent which an appear and appear and appear and a compound or agent which compound or agent which an appear and appear and appear and appear and appear and a subscript of mammalian KSR protein; stimulating the expression of mammalian KSR protein; stimulating compound or agent which and a subscript of mammalian KSR protein of a vector expression of KSR, in the presence and absence of a candidate compound or agent which activates the expression of KSR in the presence of the candidate compound or agent which activate and decrease in the expression of KSR mammalian KSR protein or a vector expression of KSR. In the presence of the candidate of the candidate compound or agent which appears KSR mines presence of the candidate of the candid Sequence 120 BP; 31 A; 36 C; (C3) comprising an expression vector and a carrier or diluent, where the expression vector is capable of expressing nucleic acid which is substantially complementary to the coding sequence of KSR RNA, or its portion/fragment, where the nucleic acid inhibits the expression of KSR; radiotherapeutic agent and an oligonucleotide which is targeted to a mRNA encoding mammalian KSR and which inhibits KSR expression; a composition diluent; a composition (C2) expression of KSR; a pharmaceutical composition (PC1) comprising (I) and a carrier or diluent; a composition (C1) comprising (I) and a carrier or substantially complementary to the CA1 region of the coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide inhibits the ovarian cancer, The present sequ **sequence** stomach cancer, esophageal cance uence represents a KSR antisense comprising one or more, chemotherapeutic or 32 G; 21 T; 0 U; 0 Other; cancer and oligonucleotide

Gaps

AEA46558
ID AEA4
XX
AC AEA4
XX
DT 25-4 RESULT 7 ₽ S Matches Query Match Best Local AEA46558 standard; DNA; 121 91 Similarity CGGACCCTAGAGGCAAAG 18 Conservative 100.0%; Score 18; DB 100.0%; Pred. No. 22; ₽P 108 0; Mismatches DB 14; 0 Length 120; Indels <u>,</u>

25-AUG-2005

(first entry)

AEA46558;

Kinase suppressor of Ras; KSR; cytostatic; neoplasm; pancreas tumor; liver tumor; skin tumor; thyroid tumor; lung tumor; urinary tract tumo gastrointestinal disease; respiratory disease; genitourinary disease; endocrine disease; bladder tumor; colon tumor; intestine tumor; head & neck tumor; leukemia; hematological disease; lymphoma; immune disorder; esophagus tumor; breast tumor; stomach tumor; nervous system tumor; neurological disease; ovary tumor; neurological disease; ovary tumor; gynecology and obstetrics; ss; angiogenesis disorder; antiangiogenic; cardiovascular disease; antisense oligonucleotide; antisense therapy; Synthetic. prostate tumor; Kinase suppressor sapiens andrology; cancer; hyperproliferation. /product= "Kinase suppressor of Ras (KSR) fragment" /note= "No start or stop codon shown" Location/Qualifiers partial 'tag= of. Ras (KSR) antisense oligonucleotide, þ lung tumor; urinary tract tumor; SEQ ID NO:25

WO2005056756-A2

03-DEC-2004; 2004WO-US040506

03-DEC-2003; 2003US-00727358

(SLOK ) SLOAN KETTERING INST CANCER RES.

R. Xing

WPI; 2005-445165/45

New oligonucleotide inhibiting kinase suppressor of Ras (KSR) euseful for inhibiting expression of mammalian KSR and treating hyperproliferative conditions such as lung cancer, skin cancer, expression, and

Claim 6; SEQ ID NO 25; 186pp; English

CC to the CA1 region of KSR; an antisense oligonucleotide (IV) comprising a CC sequence substantially complementary to nucleotides 124-243 of the coding CC sequence of mouse KSR or nucleotides 97-216 of human KSR or its portion; CC ARA6539, ARA6540 and ARA6541, and; a recombinant DNA molecule (VI) CC comprising a nucleic acid sequence which encodes on transcription an CC entisense RNA complementary to mammalian KSR RNA or its portion; a cell (VII) transfected with (VI); an expression vector (VI) capable of CC expressing a nucleic acid which is substantially complementary to the CC oligonucleotide/nucleic acid which is substantially complementary to the CC expression vector (V2) capable of Expression vector (V2) capable of Expression of KSR; an antiponucleotide which is capable of expression of KSR; and CC expression vector (V2) capable of expression of KSR; and CC expression of KSR; a pharmaceutical composition (PC1) comprising (I) and a carrier or diluent; a composition (C1) comprising (I) and a carrier or diluent; a composition one composition which is targeted to a mRNA composition (C2) comprising one or more chemotherapeutic or a mRNA composition (C2) comprising one or more chemotherapeutic or a mRNA composition (C2) comprising one or more chemotherapeutic or a mRNA composition (C2) comprising one or more chemotherapeutic or a mRNA composition (C2) comprising one or more chemotherapeutic or a mRNA composition (C2) comprising one or more chemotherapeutic or a mRNA composition (C2) comprising one or more chemotherapeutic or composition (C2) comprising composition (C2) comprising (C The invention relates to an oligonucleotide (I) which is substantially complementary to a region of kinase suppressor of Ras (KSR) RNA, where the oligonucleotide inhibits the expression of KSR. Also included are the following: an oligonucleotide (II) which is substantially complementary to a translation initiation site, 5' untranslated region, coding region or 3' untranslated region of mRNA encoding mammalian KSR; an antisense oligonucleotide (III) comprising a sequence substantially complementary radiotherapeutic agent encoding mammalian KSR (C2) comprising one or more chemotherapeutic or and an oligonucleotide which is targeted to a mRNA and which inhibits KSR expression; a composition

RESULT 8
AEA46557
ID AEA4
XX
AC AEAA
XX
DT 25-4
XX
DT 25-4
XX
EMW gene
KW gene
KW urin
KW gene

AEA46557;

AEA46557 standard; cDNA; 2601

Human kinase

suppressor

of Ras

**ب** 

(KSR1) cDNA.

(first

pancreas tumor;

urinary tract

gene; Kinase suppressor of Ras; KSR; cytostatic; neoplasm;

r; liver tumor; skin tumor; thyroid tumor; lutumor; gastrointestinal disease; respiratory disease; endocrine disease; bladder tumor; c

lung tumor;
ory disease;

colon

cc inhibiting (MI) the expression of mammalian KSR comprising contacting a colls which express KSR with a nucleic acid which is complementary to a comportating of the mRNA encoding KSR; treating or preventing (M2) a compound or agent which inhibits the expression of gf-Ras or cheightened expression of Ras in a mammal comprising administering to the caseociated with the expression of mammalian KSR compression of mammalian KSR complementary to a portion of the mRNA encoding KSR; treating or preventing (M3) a hyperproliferative condition of associated with the expression of gf-Ras or heightened expression of Ras complementary to a portion of the mRNA encoding KSR; treating to the complementary to a portion of the mRNA encoding KSR; treating or complementary to a portion of the mRNA encoding KSR; treating or expression of mammal a compound or agent which inhibits the expression of mammal a compound or agent which inhibits the expression of mammal and the procession of mammal a compound or agent which inhibits the expression of mammal and the procession of mammal and the procession of mammal and the presence of the capture of the procession of mammal and the presence of mammal and the presence of mammal and the presence and absence of a candidate compound or agent which hibits the expression of mammal and the presence and absence of a candidate compound or agent, and detecting or measuring the expression of KSR, in the presence and absence of a candidate compound or agent which hibits the expression of KSR in the presence of the candidate compound or agent inhibiting and the expression of KSR mammalian KSR in the presence of the candidate compound or agent versus in the absence of the candidate compound or agent inhibiting the expression of KSR in the presence of the candidate compound or agent which express KSR in the presence of the candidate compound or agent inhibiting and the presence of the candidate compound or agent inhibiting the expression of KSR in the presence of the candidate compound or agent which exp 밁 Ś Matches Query Match Best Local hyperproliferative condition associated with expression of gf-Ras or heightened expression of Ras in a mammal and for treating or inhibiting the progression of cancer in a mammal The cancer is chosen from pancreatic cancer, lung cancer, skin cancer, urinary tract cancer, bludder cancer, liver cancer, thyroid cancer, colon cancer, intestinal bladder cancer, liver cancer, thyroid cancer, colon cancer, breast cancer, leukemia, lymphoma, neuroblastoma, head and neck cancer, breast cancer, ovarian cancer, stomach cancer, esophageal cancer and prostate cancer. The present sequence represents a KSR antisense oligonucleotide substantially complementary to the coding sequence of KSR RNA, or in portion/fragment, where the nucleic acid inhibits the expression of (C3) comprising an expression vector and a carrier or diluent, expression vector is capable of expressing nucleic acid which 91 18; Similarity 121 CGGACCCTAGAGGCAAAG CGGACCCTAGAGGCAAAG 18 29 A; 40 100.0%; Score 18; 100.0%; Pred. No. Ç 108 0; 32 G; Mismatches 20 T; 0 DB 14; U; 0 ٥, Other; Length 121; oligonucleotide 0 inhibiting where the Gaps 0

intestine tumor; head & neck tumor; leukemia; hematological disease; lymphoma; immune disorder; esophagus tumor; breast tumor; stomach tumor; nervous system tumor; neurological disease; ovary tumor; gynecology and obstetrics; ss; angiogenesis disorder; antiangiogenic; cardiovascular disease; antisense therapy; prostate tumor; andrology; cancer; hyperproliferation.

Homo sapiens

WO2005056756-A2

03-DEC-2004; 2004WO-US040506

03-DEC-2003; 2003US-00727358

(SLOK ) SLOAN KETTERING INST CANCER RES

Kolesncik RN, Xing HR;

WPI; 2005-445165/45

New oligonucleotide inhibiting kinase suppressor of Ras (KSR) expression, useful for inhibiting expression of mammalian KSR and treating hyperproliferative conditions such as lung cancer, skin cancer, and

Example 5; SEQ ID NO 24; 186pp; English

ic portion/fragment, where the nucleic acid innibits the expression of mammalian KSR comprising contacting calls which express KSR with a nucleic acid which is complementary to a portion of the mRNA encoding KSR; treating or preventing (M2) a comper condition associated with the expression of gf-Ras or the heightened expression of Ras in a mammal comprising administering to the mammal a compound or agent which inhibits the expression of mammalian KSR compound or agent which inhibits the expression of mammalian KSR compound or agent which inhibits the expression of Ras in a mammal comprising expression of gf-Ras or heightened expression of Ras in a mammal comprising expression of gf-Ras or heightened expression of Ras compound therapeutically effective amount of a nucleic acid which is complementary to a portion of the mRNA encoding KSR; treating or inhibiting (M4) the progression of cancer in a mammal comprising administering to a mammal a compound or agent which inhibits the The invention relates to an oligonucleotide (I) which is substantially CC complementary to a region of kinase suppressor of Ras (KSR) RNA, where the oligonucleotide inhibits the expression of KSR. Also included are the CC following: an oligonucleotide (II) which is substantially complementary CC to a translation initiation site, 5' untranslated region, coding region or 3' untranslated region of mRNA encoding mammalian KSR; an antisense oligonucleotide (III) comprising a sequence substantially complementary to the CA1 region of KSR; an antisense oligonucleotide (IV) complementary to the CA1 region of KSR; an antisense oligonucleotide (IV) comprising a sequence substantially complementary to mammalian KSR or its portion; an antisense oligonucleotide (V) comprising a sequence chosen from CC AEA46539, AEA46540 and AEA46541, and; a recombinant DNA molecule (VI) comprising a nucleic acid sequence which encodes on transcription an CC expressing a nucleic acid which is substantially complementary to the coding sequence of KSR RNA, or its portion/fragment, where the cligonucleotide which is substantially complementary to the coding sequence of KSR RNA, or its portion/fragment, where the cligonucleotide which is substantially complementary to the CA1 region of the coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide which is substantially complementary to the CA1 region of the coding sequence of CC expression vector (V2) capable of expression of KSR; an antisense of KSR RNA, or its portion/fragment, where the oligonucleotide which is targeted to a mRNA contoberapeutic agent and an oligonucleotide which is targeted to a mRNA contoberapeutic open that a carrier or contacting an expression vector and a carrier or diluent, where the nucleic acid which is targeted to a mRNA contoberapeutic open than the nucleic acid which is complementary to the coding sequence of KSR RNA, or its portion of KSR

> CC the expression of mammalian KSR protein or a vector expressing mammalian CC KSR; identifying compounds or agents which inhibit the expressing of KSR, CC paiddate compound or agent, and detecting or measuring the expression of KSR CC candidate compound or agent, and detecting or measuring the expression of CC KSR in the presence and absence of a candidate compound or agent, where a CC compound or agent versus in the absence of the candidate compound or agent indicates that the compound or agent inhibits the expression of KSR in the presence of the candidate compound or agent indicates that the compound or agent inhibits the expression of KSR compound or agent inhibits the expression of KSR comprising contacting cells which express KSR with (I) is useful for inhibiting the conferring radiosensitivity to ionizing radiation in tumor cells in a mammal. (I) Is useful for inhibiting angiogenesis in a mammal. (I) Is useful for inhibiting angiogenesis in a mammal. The methods are useful for treating or preventing the progression of Ras in a mammal and for treating or inhibiting the progression of grancer, lung cancer, skin cancer, urinary tract cancer, cancer, lung cancer, thyroid cancer, colon cancer, intestinal CC cancer, leukenia, lymphoma, neuroblastoma, head and neck cancer, intestinal CC cancer, ovarian cancer, stomach cancer, esophageal cancer and prostate cancer. The present sequence represents human KSRI cDNA. angiogenesis in a mammal comprising administering to a mammal a therapeutically effective amount of a compound or agent which activates expression of mammalian KSR protein; inhibiting angiogenesis of tumor cells in a mammal, by administering to a mammal a compound or agent whic inhibits the expression of mammalian KSR protein; inhibiting or reducing VEGF expression or activity in a mammal, by administering a compound or agent that inhibits the expression of mammalian KSR protein; stimulating or reducing which

Sequence 2601 BP; 565 A; 818 C; 772 G; 446 T; 0 U; 0 Other;

Matches Query Match Best Local 18; Similarity Conservative 100.0%; 0, Score 18; Pred. No. Mismatches 25 BB 14; <u>,</u> Length 2601; Indels 0, Gaps 0,

밁 S 187 CGGACCCTAGAGGCAAAG 204 CGGACCCTAGAGGCAAAG 18

AEA46544 standard; cDNA; 4034 BP

AEA46544

25-AUG-2005 (first entry)

Mouse kinase suppressor of Ras (KSR) cDNA

RESULT 9
AEA46544
ID AEA4
XX AEA4
XX AEA4
XX AEA4
XX AEA4
XX Mous
XX Mous
XX Mous
XX Gene
KW Panc
KW Panc
KW Indry
KW Indry
KW Geni
KW Indry
KW Geni
KW Card
KW Card
XX Mus
XX WO20
XX WO20
XX WO20
XX WO20 gene; Kinase suppressor of Ras; KSR; cytostatic; neoplasm; pancreas tumor; liver tumor; skin tumor; thyroid tumor; lumor; urinary tract tumor; gastrointestinal disease; respiratory disease; genitourinary disease; endocrine disease; bladder tumor; colon tumor; intestine tumor; head & neck tumor; leukemia; hematological disease; lymphoma; immune disorder; esophagus tumor; breast tumor; stomach tumor; nervous system tumor; neurological disease; ovary tumor; gynecology and obstetrics; ss; angiogenesis disorder; antiangiogenic; cardiovascular disease; antisense therapy; prostate tumor; andrology; cancer; hyperproliferation

ďB

Location/Qualifiers 83. .2704

product= "Mouse kinase suppressor of Ras (KSR)"

WO2005056756-A2

23-JUN-2005

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CC expression vector (V2) capable of expressing an oligonuclectide which is capable of expressing an oligonuclectide which is CC substantially complementary to the CA1 region of the coding sequence of CC KSR RNA, or its portion/iragment, where the oligonuclectide inhibits the capable of expression of KSR; a pharmaceutical composition (PC1) comprising (I) and carrier or diluent; a composition (C2) comprising one or more chemotherapeutic or CC diluent; a composition (C2) comprising one or more chemotherapeutic or cardiotherapeutic agent and an oligonucleotide which is targeted to a mRNA CC encoding mammalian KSR and which inhibits KSR expression; a composition (C2) comprising an expression vector and a carrier or diluent, where the coding sequence of KSR RNA, or its gortion of transpent, where the nucleic acid inhibits the expression of KSR; transpention of KSR comprising contacting (C2) comprising (M1) the expression of mammalian KSR comprising contacting (C2) comprising (M1) the expression of mammal comprising doministering to a portion of the mRNA encoding KSR; treating or preventing (M2) a hyperproliferative condition associated with the expression of gf. Ras or heightened expression of Ras in a mammal comprising administering to the mammal therapeutically effective amount of a nucleic acid which is complementary to a portion of the mRNA encoding KSR; treating or capable or expression of tumor (C2) inhibits the expression of mammalian KSR protein; inhibiting angiogenesis of tumor (C3) expression of activity in a mammal, by administering to the mammal comprising and protein; inhibits the expression of KSR. (and in the protein of mammalian KSR protein, inhibits the expression of tumor or agent which inhibits the expression of KSR. (and in a mammal) a compound or agent which activates the expression of mammalian KSR protein, inhibiting a compound or the expression of KSR. (and in a mammal) and compound or agent which inhibits the expression of KSR. (and in a mammal) and compound or agent which inhibits the expre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to a translation initiation site, 5' untranslated region, coding region or 3' untranslated region of mRNA encoding mammalian KSR; an antisense oligonucleotide (III) comprising a sequence substantially complementary to the CA1 region of KSR; an antisense oligonucleotide (IV) comprising a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expressing a nucleic acid which is substantially complementary to the coding sequence of KSR RNA, or its portion/fragment, where the oligonucleotide/nucleic acid inhibits the expression of KSR; an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEA46539, AEA46540 and AEA46541, and, a recombinant DNA molecule (VI) comprising a nucleic acid sequence which encodes on transcription an antisense RNA complementary to mammalian KSR RNA or its portion; a cell (VII) transfected with (VI); an expression vector (VI) capable of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an oligonucleotide (I) which is substantially complementary to a region of kinase suppressor of Ras (KSR) RNA, where the oligonucleotide inhibits the expression of KSR. Also included are the following: an oligonucleotide (II) which is substantially complementary
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identifying compounds or agents which inhibit the expression of KSR, ncubating a cell expressing KSR in the presence and absence of a idate compound or agent, and detecting or measuring the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2005-445165/45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and
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Ś CC KSR in the presence and absence of a candidate compound or agent, where a CC decrease in the expression of KSR in the presence of the candidate CC compound or agent versus in the absence of the candidate compound or agent indicates that the compound or agent inhibits the expression of KSR compound or agent inhibits the expression of KSR CC expression of mammalian KSR compound in the candidate compound or gent inhibits the expression of KSR CC with (1), where expression of mammalian KSR is inhibited. (1) Is useful for conferring radiosensitivity to ionizing radiation in tumor cells in a CC mammal. (1) Is useful for inhibiting angiogenesis in a mammal. (1) Is useful for mammal. (2) Is useful for inhibiting angiogenesis in a mammal. (3) Is useful for inhibiting angiogenesis in a mammal. (4) Is useful for treating or preventing common com 밁 RESULT 10 Query Match .Best Local S Matches 18 Isolated kinase suppressor of ras protein - used to identify lead compounds and in diagnosis or treatment of disease associated with 13-DEC-1995; WO9721820-A2 Cus Mouse Ksr-1 (kinase suppressor of Ras) cDNA. 13-SEP-1997 AAT72155; AAT72155 standard; cDNA; 4094 BP Sequence 4034 BP; Rubin G, 13-DEC-1996; 19-JUN-1997 Mus musculus Kinase (REGC ) UNIV CALIFORNIA 1997-332791/30 se suppressor of Ras; Ksr-1; signal transduction; cell growth; differentiation; gene therapy; diagnosis; ds. 296 CGGACCCTAGAGGCAAAG 313 18; Similarity AAW19918 CGGACCCTAGAGGCAAAG 18 ဝူ Therrien M, Ksr signal Conservative (first entry) 95US-00571758. 96WO-US019941. Location/Qualifiers 83. .2704 /\*tag= a 906 A; 1200 C; 100.0%; 100.0%; Score 18; 100.0%; Pred. No. Chang 0, Η, Mismatches 1171 G; Karim F, 757 T; 0 U; 0 Other; DB 14; Wassarman 0 Length 4034; Indels associated with Ksr 0 Gaps 0,

A CDNA clone (AAT72155) includes a complete open reading frame that codes for mouse kinase suppressor of Ras (AAW19918) designated Kgr-1, a novel

49pp;

English

SSSSSSSSSS

0,

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Query Match
Best Local S
Matches 18
                                             The invention comprises antisense oligonucleotides which are designed to inhibit the expression of the kinase suppressor of Ras (KSR) gene. The antisense oligonucleotides of the invention are useful for treating or inhibiting the progression of cancer, such as: pancreatic cancer, lung cancer, skin cancer, urihary tract cancer, bladder cancer, liver cancer, thyroid cancer, colon cancer, intestinal cancer, breast cancer, ovarian cancer, stomach cancer, head and neck cancer, oesophageal cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein kinase involved in the regulation of cell growth and differentiation. It was isolated from a mouse PCC4 teratocarcinoma cell line with a probe corresponding to hb, a sequence that had been identified as showing similarity to Drosophila melanogaster Ksr (see als AAT72153). Drosophila virilis and human Ksr sequences (see also AAT72154 AAT82156-57) have also been identified. These Ksr homologues define a novel class of kinases related to raf kinases. Ksr mucleic acids can be used to produce Ksr polypeptides, as probes to identify mutant Ksr alleles associated with disease, and in gene therapy applications
                                                                                                                                                                                                                                                                          Example 1;
                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotides complementary to a region of kinase suppressor of Ras (KS) RNA which inhibit KSR expression, useful for treating or inhibiting progression of cancer, e.g. pancreatic, lung, skin, or bladder cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-2002; 2002US-0384228P
03-APR-2003; 2003US-0460023P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse kinase
                      prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kolesnick RN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003101386-A2
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2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   intestinal cancer; breast cancer; ovarian cancer; stomach cancer;
head cancer; neck cancer; oesophageal cancer; prostate cancer; le
lymphoma; neuroblastoma; mouse; murine; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pancreatic cancer; lung cancer; skin cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG75394 standard; cDNA; 4094 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAY-2003; 2003WO-US016961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bladder cancer; liver cancer; thyroid cancer; colon cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4094 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SLOK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2004-053335/05.
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cancer,
encodes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CGGACCCTAGAGGCAAAG 18
                                                                                                                                                                                                                                                                       SEQ ID NO 11; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                suppressor of Ras (KSR) coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KETTERING INST CANCER RES.
leukaemia, ly
the mouse KSR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      924 A; 1212 C; 1193 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                         lymphoma, and neuroblastoma. The present cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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The invention relates to an oligonuclectide (I) which is substantially complementary to a region of kinase suppressor of Ras (KSR) RNA, where CC the oligonuclectide inhibits the expression of KSR. Also included are the Cf following: an oligonuclectide (II) which is substantially complementary to a translation initiation site, 5' untranslated region, coding region CC or 3' untranslated region of mRNA encoding mammalian KSR; an antisense CC oligonucleotide (III) comprising a sequence substantially complementary to the CA1 region of KSR; an antisense oligonucleotide (IV) comprising a sequence substantially complementary to nucleotides 124-243 of the coding CC sequence of mouse KSR or nucleotides 97-216 of human KSR or its portion; CC an antisense oligonucleotide (V) comprising a sequence chosen from CC AEA46539, AEA46540 and AEA46541, and; a recombinant DNA molecule (VI) comprising a nucleic acid sequence which encodes on transcription an CC comprising a nucleic acid sequence which encodes on transcription an antisense RNA complementary to mammalian KSR RNA or its portion; a cell (VII) transfected with (VI); an expression vector (VI) capable of expressing a nucleic acid which is substantially complementary to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ÖS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New oligonucleotide inhibiting kinase suppressor of Ras (KSR) expression, useful for inhibiting expression of mammalian KSR and treating hyperproliferative conditions such as lung cancer, skin cancer, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kinase suppressor of Ras; KSR; cytostatic; neoplasm; pancreas invertumor; skin tumor; thyroid tumor; lung tumor; urinary trigistrointestinal disease; respiratory disease; genitourinary disease; separatorine disease; respiratory disease; genitourinary disease; lisease; lumor; intestine tumor; endocrine disease; bladder tumor; colon tumor; intestine tumor; head & neck tumor; leukemia; hematological disease; lymphoma; immune disease; esophagus tumor; breast tumor; stomach tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 8; SEQ ID NO 8; 186pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kolesncik RN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-DEC-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gymecology and obstetrics; ss; angiogenesis disorder; antiangiogenic; cardiovascular disease; antisense oligonucleotide; antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4094 BP; 924 A; 1212 C; 1193 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nervous system tumor; neurological disease; ovary tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kinase suppressor of Ras (KSR) antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-AUG-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 CGGACCCTAGAGGCAAAG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2003US-00727358
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 4094;
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cytostatic; stomach; Thyroid cancer related gene sequence SEQ ID NO:5825

cancer; colon; breast; ovary; oesophagus; kidney; thyroid;

ng; prostate; pancreas; carcinoma; antitumour; cancerous; gene therapy; antineoplastic; Wilm's tumour; adenocarcinoma;

15-MAY-2002 ABL67488;

(first entry)

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CC angiogenests in a mammal comprising administrating the expression of mammal and the expression of a mammal and the expression of a compound or agent which activates CC the expression of mammalian KSR protein or a vector expressing mammalian CC KSR; identifying compounds or agents which inhibit the expression of KSR, CC by incubating a cell expressing KSR in the presence and absence of a candidate compound or agent, and detecting or measuring the expression of KSR. CC candidate compound or agent, and detecting or measuring the expression of KSR in the presence of the candidate compound or agent versus in the absence of the candidate compound or agent indicates that the compound or agent inhibiting the expression of KSR in the presence of the candidate compound or agent indicates that the compound or agent inhibiting the capturession of mammalian KSR comprising contacting cells which expression of KSR with (1), where expression of mammalian KSR is inhibited. (1) Is useful for conferring radiosensitivity to ionizing radiation in tumor cells in a CC mammal. (1) Is useful for inhibiting angiogenesis in a mammal. (1) Is useful for inhibiting or reducing VEGF expression or activity in a CC useful for inhibiting or reducing VEGF expression of gf-Ras or the progression of cancer in a mammal and for treating or inhibiting CC the progression of Ras in a mammal and for treating or inhibiting CC the progression of cancer, skin cancer, urinary tract cancer, breast CC cancer, leverent all prophoma, neuroblastoma, head and neck cancer, breast CC cancer, orarian cancer, stancer, esonbaceal cancer and prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC (C3) comprising an expression vector and a carrier or diluent, where the CC expression vector is capable of expressing nucleic acid which is cubstantially complementary to the coding sequence of KSR RNA, or its portion/fragment, where the nucleic acid inhibits the expression of KSR; Cinhibiting (M1) the expression of mammalian KSR comprising contacting (C1) inhibiting (M2) the expression of mammalian KSR complementary to a contion of the mRNA encoding KSR; treating or preventing (M2) a portion of the mRNA encoding KSR; treating or preventing (M2) a hightened expression of Ras in a mammal comprising administering to the mammal a compound or agent which inhibits the expression of mammalian KSR protein; treating or preventing (M3) a hyperproliferative condition associated with the expression of Ras in a mammal or administering to the mammal therapeutically effective amount of a nucleic acid which is complementary to a portion of the mRNA encoding KSR; treating or inhibiting M3) the progression of the mRNA encoding KSR; treating or
                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inhibiting (M4) the progression of cancer in a mammal comprising administering to a mammal a compound or agent which inhibits the administering to a mammal an KSR protein; inhibiting angiogenesis of tumor expression of mammalian KSR protein; to a mammal a compound or agent which calls in a mammal, by administering to a mammal a compound or reducing thibits the expression of mammalian KSR protein; inhibiting a compound or agent that inhibits the expression of mammal, by administering a compound or agent that inhibits the expression of mammalian KSR protein; simulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonuclectide/nucleic acid inhibits the expression of KSR; an expression vector (V2) capable of expressing an oligonuclectide which is substantially complementary to the CA1 region of the coding sequence of KSR RNA, or its portion/fragment, where the oligonuclectide inhibits the expression of KSR; a pharmaceutical composition (PC1) comprising (I) and a carrier or diluent; a composition (C1) comprising (I) and a carrier or diluent; a composition (C2) comprising one or more chemotherapeutic or radiotherapeutic agent and an oligonuclectide which is targeted to a mRNA encoding mammalian KSR and which inhibits KSR expression; a composition
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 16 BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               coding sequence of KSR RNA,
                                                                                                                                                                                                                                                  Local
16
                                                                                                                                                                                                    16;
                                                                                                  w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ovarian cancer, stomach cancer, esophageal cancer and prostate The present sequence represents a KSR antisense oligonucleotid
                                                                                                                                                                                                                                                  Similarity
                                                                                                  GACCCTAGAGGCAAAG 18
                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  Þ
                                                                                                                                                                                                                                                                                                                                                                                             5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or its portion/fragment, where the
                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                  Score 16;
Pred. No.
                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                  DB 14;
. 2.2e+02;
                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                Length 16;
                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oligonucleotide
                                                                                                                                                                                                    0;
                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                         0
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RESULT 13 ABL67488/C ID ABL67488 standard; DNA; 376 BP XX

Screening for anti-neoplastic agent involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, and determining a change

determining a change

tested for anti-neoplastic activity, and on of a gene of a signature gene set.

gene of a

expression

2002-188264/24.

Carter

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02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
03-OCT-2000
03-OCT-2000
03-OCT-2000
                                                           03-OCT-2000;
01-NOV-2000;
01-NOV-2000;
                                                                                                                                                                                                                                                                                  30-MAY-2001; 2001WO-US010838
                                                                                                                                                                                                                                                                                            13-DEC-2001
                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                            29-SEP-
                                                                                                                                      28-SEP-
                                                                                                                                           28-SEP-2000;
                                                                                                                                                28-SEP-2000;
                                                                                                                                                     28-SEP-2000;
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                                                                                                                                                               28-SEP-2000;
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                                                                                                                                                                                                                                                     20-SEP-2000;
                                                                                                                                                                                                                                                              05-JUN-2000;
18-SEP-2000;
                                                                                                                                                                                                                                                                        05-JUN-2000;
                                                                                                                                                                               27-SEP-2000;
                                                                                                                                                                                    27-SEP-2000;
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                                                                                                                                                                                                                                      22-SEP-2000;
                                                                                                                                                                                                                                                          L8-SEP-2000;
                                                 (AVAL-)
                                                  AVALON
                                  Augustus M, Weaver Z;
                                                                2000US-0237606P.
2000US-0237608P.
2000US-0244867P.
                                                                                                                                           2000US-0236109P
                                                                                                                                                                                    2000US-0235711P
                                                                                                                                                                                              2000US-0235637P
                                                                                                                                                                                                         2000US-0235134P
                                                                                                                                                                                                              2000US-0235082P
                                                 PHARM.
                                        č,
                                        Ebner
                                        Ŗ,
                                        Endress
                                        ຸດ
                                        Horrigan
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RESULT 14
ABL69201/C
ID
ABL69201 standard; DNA; 376 BE
XX
AC
ABL69201;
XX
DT
15-MAY-2002 (first entry)
XX
DE
Prostate cancer related gene s
XX
Human; cancer; colon; breast;
XW
stomach; lung; prostate; panc;
XW
Gytostatic; gene therapy; anti
XW
gene; ds.
XX
OS
Homo sapiens.
XX
PD
13-DEC-2001, 2001WO-US010838.
XX
PF
30-MAY-2001; 2000US-0209473P
PR
05-JUN-2000; 2000US-0239613P
PR
18-SEP-2000; 2000US-023313P
PR
18-SEP-2000; 2000US-0234034P
PR
20-SEP-2000; 2000US-0234094P
PR
20-SEP-2000; 2000US-023450P
PR
20-SEP-2000; 2000US-023450P
PR
22-SEP-2000; 2000US-0234924P
PR
23-SEP-2000; 2000US-0234924P
PR
25-SEP-2000; 2000US-023563P
PR
25-SEP-2000; 2000US-023563P
PR
25-SEP-2000; 2000US-023563P
PR
26-SEP-2000; 2000US-023563P
PR
27-SEP-2000; 2000US-023563P
PR
27-SEP-2000; 2000US-023563P
PR
27-SEP-2000; 2000US-023563P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC neoplastic agent. The method involves exposing cells to a colon, breast, and the data is sufficient to the data of the data colon, breast, stomach, lung, thyroid, oversite a colon, breast, stomach, lung, thyroid, oversite a colon, breast, stomach, lung, thyroid, oversite are finitization, and the data is sufficient to convey the chemical structure and/or corporation of the agent, and can be used for producting a product which is convey the chemical structure and/or properties of the agent, and can be used for producting a product which is convey the chemical structure and/or properties of the agent. MI can be used in the treatment of cancer such compostries of the agent, and can be used for producting a product which is colon, breast, stomach, lung, thyroid, oesophageal, ovarian, kidney, cascinoma, rescinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ঠ
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 5825; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   107 GGACCCTAGAGGCAGAG 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ν
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGACCCTAGAGGCAAAG 18
                                                                                2000US-0235077P.
2000US-0235082P.
2000US-0235134P.
2000US-0235280P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP;
                          2000US-0235638P.
2000US-0235638P.
2000US-0235711P.
                                                                                                                                                                               2000US-0234567P.
2000US-0234923P.
                                                                                                                                                                                                                                         2000US-0234009P.
2000US-0234034P.
2000US-0234052P.
                                                                                                                                                                                                                                                                                                 2000US-0209531P.
2000US-0233133P.
2000US-0233617P.
                                                                                                                                                                                                                       2000US-0234509P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         colon; breast; ovary; oesophagus; kidney; thyroid;
prostate; pancreas; carcinoma; antitumour; cancerous;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 related gene sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            78 A; 132 C; 76 G;
                                                                                                                                                              -0234924P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 85.6%;
94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      describes a method (MI) for screening for an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antineoplastic; Wilm's tumour; adenocarcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            구.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ID NO:7538.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 376;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 15
ADF80786/c
ID ADF807
XX
AC ADF807
XX
AC ADF807
XX
DT 26-FEB

ADF80786

standard; DNA; 388

26-FEB-2004 ADF80786;

(first entry)

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107

91

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δ
                                                                                     cc neoplastic agent. The method involves exposing cells to a Chemical agent cobe tested for anti-neoplastic activity, determining a change in cc expression of at least one gene (I) of a signature gene set, where (I) cc comprises a sequence (S) selected from 8447 sequences (given in ABL61664 to ABL70110), or is at least 95% identical to (S), where a change in expression is indicative of anti-neoplastic activity. (I) has cytostatic cativity and can be used in gene therapy. MI can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a result of MI, and the data is sufficient to convey the chemical structure and/or properties of the agent. MI can be used in the treatment of cancer such as colon, breast, stomach, lung, thyroid, oesophageal, ovarian, kidney, prostate or pancreatic cancer, adenocarcinoma, carcinoma, clear cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's cumour
                             Matches
                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-SEP-2000;
02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
03-OCT-2000;
03-OCT-2000;
03-OCT-2000;
03-OCT-2000;
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27-SEP-2000;
28-SEP-2000;
28-SEP-2000;
                                                                                     Sequence 376 BP;
                                                                                                                                                                                                                                                                                                            The present invention describes a method (M1) for screening for an neoplastic agent. The method involves exposing cells to a chemical to be tested for anti-neoplastic activity, determining a change in
                                                                                                                                                                                                                                                                                                                                                                                                Screening for anti-neoplastic agent involves exposing cells to a agent to be tested for anti-neoplastic activity, and determining in expression of a gene of a signature gene set.
                                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-188264/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Soppet
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29-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Young
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28-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AVAL-) AVALON PHARM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PE,
                           16;
 2 GGACCCTAGAGGCAAAG 18
                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 7538; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Augustus M,
Weaver Z;
                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0237278P.
2000US-0237294P.
2000US-0237295P.
2000US-0237316P.
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2000US-0235863P.
2000US-0236028P.
2000US-0236032P.
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2000US-0237598P.
2000US-0237604P.
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2000US-0244867P
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2000US-0237172P.
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2000US-0236842P.
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2000US-0236034P.
                                                                                     78
                                            85.6%;
94.1%;
                                                                                   A; 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Carter KC,
                                                                                     Ç
                              0
                                            Score 15.4; DB Pred. No. 5.1e+(
                                                                                     76 G;
                              Mismatches
                                                                                     90
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                                                                                     H
                                            5.1e+02;
                                                                                     0 U;
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                                                                                      0
                                                       Length 376;
                                                                                      Other;
                              Indels
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                             Gaps
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a change
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RESULT 16
AAK88398/c
ID AAK883
XX AAK883
AC AAK883
XX O5-NOV
XX Human
XX Human;
KW Human;
KW digest
XX digest
XX WO2001
XX PP 02-AUC
XX 17-JAN
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                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 16
                                                                           Human; digestive system antigen; gene therapy; cancer; appendicitis; ulcerative colitis; infection; Hirschsprung's disease; chronic colitis; digestive system disorder; Meckel's diverticulum; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eils
                                                                                                                                                                                                                                                                                                                            The present invention relates to a method (M1) for determining the subtype of leukaemia cells and whether a patient sample contains leukaemia cells. The method comprises determining the expression profile of a group of markers in a patient sample. The method is useful for determining the presence of leukaemia cells, its types or subtypes, and for the preparation of a medicament for treating leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                               Determining the subtype of leukemia cells and whether a patient sample contains leukemia cells or other cells, useful for treating leukemia, comprises determining the expression profile of a group of markers in a
 17-JAN-2001; 2001WO-US001324.
                                                                                                                    Human digestive
                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 1342; 2938pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-505037/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Haferlach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (DEKR-)
(UYLU-)
(HAFE/)
(SCHO/)
                    02-AUG-2001.
                                       WO200155314-A2
                                                          Homo sapiens.
                                                                                                                                        05-NOV-2001
                                                                                                                                                           AAK88398;
                                                                                                                                                                              AAK88398 standard; cDNA; 514 BP
                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                      patient sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-NOV-2001;
30-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-NOV-2002; 2002WO-EP012303.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003039443-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytostatic; Gene therapy; leukaemia; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Leukaemia-related DNA sequence #1342.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KERN/)
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                                                                                                                                                                                                                               360
                                                                                                                                                                                                                              2 GGACCCTAGAGGCAAAG 18
|||||||||||||
60 GGACCCTAGAGGCCAAG 344
                                                                                                                                                                                                                                                                     l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SCHOCH C.
KERN W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DEUT KREBSFORSCHUNGSZENTRUM
UNIV LUDWIG MAXIMILIANS.
HAFERLACH T.
                                                                                                                                                                                                                                                                                                            388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        h T, Schoch C,
Brors B, Merg
                                                                                                                                                                                                                                                                      Conservative
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2002EP-00009758.
                                                                                                                                       (first entry)
                                                                                                                   system antigen coding sequence SEQ ID
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Mergenthaler S;
                                                                                                                                                                                                                                                                              85.6%;
94.1%;
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Pred. No. 5.1e+02;
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04-FEB-2000;
24-FEB-2000;
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19-MAY-2000;
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2000US-0230437P.
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Polynucleotides encoding digestive system antigens, useful
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2000US-0251990P.
2000US-0254097P.
2001US-0259678P.
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Matches 16
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17-MAR-2000;
18-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; colon cameer; congenital abnormality; infection; colitis; inflammatory bowel disease; IBD; neoplastic disorder; gene therapy; intestinal inflammatory disorder, malabsorption syndroms; gastric; sigmoid disease; antibacterial; antiviral; antiinflammatory; cytostatic;
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                          2000US-0189874P.
2000US-0198123P.
2000US-0205515P.
2000US-0214886P.
2000US-0214886P.
2000US-0215135P.
2000US-0216847P.
2000US-0216847P.
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2000US-0217487P.
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2000US-0218293P.
2000US-0218293P.
2000US-0228513P.
2000US-0224513P.
2000US-0225261P.
2000US-0225267P.
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2000US-0180628P.
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Pred. No. 5
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17-NOV-2000; 17-NOV-2000; 17-NOV-2000; 17-NOV-2000; 17-NOV-2000;

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GENOME

S-0225270P. S-0225447P. S-0225757P.

30-AUG-2000; 01-SEP-2000; 01-SEP-2000; 01-SEP-2000; 01-SEP-2000; 01-SEP-2000;

2000US 2000US 2000US

3-0229345P.

2000US-2000US-

-0226681P

2000US

S-0228924P. S-0229287P. S-0229343P. S-0229344P.

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The present invention relates to the isolation of novel human colon cassociated polypeptides (AAU22468-AAU22701), and the CDNA and genomic sequences encoding for them. The sequences of the invention are useful in the diagnosis, treatment, prevention and/or prognosis of disorders of the colon including colon cancer, congenital abnormalities (e.g. atresia and stenosis), bacterial and viral infections, inflammatory bowel disease (CIBD), neoplastic cell disorders (e.g. polyps and adenomas, intestinal colonitariory, malabsorption syndromes (e.g. lactose incolerance), intestinal construction and sigmoid diseases. The polynucleotides sequences of the colonitarion can also be used in gene therapy. AAS39348-AAS39581 represent coff the invention. Note: The sequence data for this pattent did not form cart of the printed specification, but was obtained in electronic format colonication from MIPO at ftp.wipo.int/pub/published_pct_sequences
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10-DEC-2000;
05-DEC-2000;
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08-NOV-2000;
08-NOV-2000;
                                                                                                                                                                                                                                                                                                             Isolated polypeptide for treating, preventing and/ or prognosing disorders related to the colon including colon cancers and also for testing and detection e.g. diagnosis.
   Sequence 514
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P-PSDB; AAU22619.
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17-NOV-2000;
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    BP; 162
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    A; 115
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14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 21-SEP-2000; 21-SEP-2000; 25-SEP-2000; 25-SEP-2000;

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2000US

2000US-0235484P 2000US-0235834P 2000US-0235836P 2000US-023632P 2000US-0236367P 2000US-0236369P 12-SEP-2000; 14-SEP-2000; 14-SEP-2000;

S-0231968P. S-0232397P. S-0232398P.

-0232080P.

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2000US-0246523P.
2000US-0246527P.

Query Match 85.6%; Best Local Similarity 94.1%; Matches 16; Conservative

Score 15.4; DB 5; Pred. No. 5.1e+02; 0; Mismatches 1;

Length 514;

Indels

0;

Gaps

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 14-AUG-2000
18-AUG-2000
22-AUG-2000
22-AUG-2000
22-AUG-2000
22-AUG-2000
23-AUG-2000
30-AUG-2000
01-SEP-2000
01-SEP-2000
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05-SEP-2000
06-SEP-2000
08-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene therapy; ss; gene; cancer; liver disorder; hepatitis;
neural disorder; Alzheimer's disease; human; colon.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human novel colon related polypeptide cDNA SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-2003
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2000US-0225758P

2000US-0226279P

2000US-022688P

2000US-022688P

2000US-0227182P

2000US-0227182P

2000US-022798924P

2000US-02298924P

2000US-0229343P

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2000US-0184664P.
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2000US-01846350P.
2000US-0199076P.
2000US-0199076P.
2000US-0199076P.
2000US-0215133P.
2000US-0214886P.
2000US-021644P.
2000US-021649P.
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2000US-0225447P.
2000US-0225757P.
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01-NOV-2000;
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08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
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20-OCT-2000;
2000US-0249210P
2000US-02492110P
2000US-02492112P
2000US-02492113P
2000US-02492149
2000US-0249214P
2000US-0249215P
2000US-0249216P
2000US-0249217P
2000US-0249218P
2000US-0249218P
2000US-0249248P
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2000US-0241785P.
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2000US-0241808P.
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2000US-0246474P.
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2000US-0236370P.
2000US-0236802P.
2000US-0237037P.
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2000US-0237039P.
2000US-0237040P.
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2000US-0231413P.
2000US-0231414P.
2000US-0232080P.
2000US-0232081P.
2000US-0232081P.
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2000US-0234274P.
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2000US-0235484P.
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2000US-0233063P.
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2000US-0241826P.
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2000US-0246613P.
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2000US-0246609P.
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2000US-0240960P.
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RESULT 19
ADF82177/c
ID ADF821
XX ADF821
XX ADF821
XX Leukae
XX Cycost
XX Cycost
XX Unider
XX W0200:
XX PP 04-N0
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Best Local :
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01-DEC-2000;
05-DEC-2000;
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17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                         The invention relates to an isolated nucleic acid molecule encoding a polypeptide. The nucleic acid is useful for preparing a medicament for preventing, treating or medicating a medical condition e.g. cancer, liver disorders such as hepatitis or neural disorders such as Alzheimer's disease. The present sequence represents a human cDNA encoding a novel colon related polypeptide. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from USPTO at
                                                                                                             Leukaemia-related DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid molecule, useful for preparing a medicament for preventing, treating or ameliorating a medical condition e.g. cancer, liver disorders or neural disorders.
   04-NOV-2002; 2002WO-EP012303
                                                                                       Cytostatic;
                                              WO2003039443-A2
                                                                    Unidentified
                                                                                                                                   26-FEB-2004
                                                                                                                                                         ADF82177;
                                                                                                                                                                              ADF82177
                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                               seqdata.uspto.gov/sequence.html?DocID=20030050231.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; SEQ ID NO 162; 216pp; English.
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                                                                                                                                                                              standard;
                                                                                                                                                                                                                                    GGACCCTAGAGGCCAAG 414
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                                                                                                                                                                                                                                                                                Conservative
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2000US-0251868P.
2000US-0251869P.
                                                                                                                                                                                                                                                                                                                            BP;
                                                                                                                                  (first entry)
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2000US-0256719P.
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2000US-0250160P.
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2000US-0249297P.
2000US-0249299P.
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                                                                                                                                                                                                                                                                                                                            162
                                                                                      therapy;
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                                                                                                                                                                                                                                                                                         85.6%;
94.1%;
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                                                                                                                                                                              ВP
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                                                                                                                                                                                                                                                                                         Score 15.4; DB 9; Pred. No. 5.1e+02;
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                                                                                                              #2733
                                                                                                                                                                                                                                                                                                                            96 G;
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                                                                                                                                                                                                                                                                                                                            141 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                      Length 514;
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RESULT 20
ADQ24821/c
ID ADQ248
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Best Local :
                                                                                                                                                     10-JUN-2004.
                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                               Human soft tissue
                                                                                                                                                                                                                                                                   26-AUG-2004
                                                                                                                                                                                                                                                                                                            ADQ24821 standard; DNA; 1797 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a method (M1) for determining the subtype of leukaemia cells and whether a patient sample contains leukaemia cells. The method comprises determining the expression proof a group of markers in a patient sample. The method is useful for of a group of markers in a patient sample. The method is useful for determining the presence of leukaemia cells, its types or subtypes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 2733;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    comprises determining the expression profile of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determining the subtype of leukemia cells and whether a patient sample contains leukemia cells or other cells, useful for treating leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-505037/47.
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30-APR-2002;
                                                                    Aziz N,
                                                                                                            26-NOV-2002; 2002US-0429739P
                                                                                                                                26-NOV-2003; 2003WO-US038193
                                                                                                                                                                        WO2004048938-A2
                                                                                                                                                                                                                          soft tissue
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                                                                                         (PROT-)
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                                                  2004-441208/41.
                                                                                                                                                                                            sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DEUT KREBSFORSCHUNGSZENTRUM UNIV. LUDWIG MAXIMILIANS.
                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                preparation of a medicament
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                                                                                          PROTEIN DESIGN
                                                                   Ginsburg WM,
                                                                                                                                                                                                                                                                                                                                                             GGACCCTAGAGGCCAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                             986 BP; 269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brors
                                                                                                                                                                                                                          sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001EP-00126244
2002EP-00009758
                                                                                                                                                                                                                                                                   (first
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s B, Mergenthaler
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94.1%;
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                                                                                                                                                                                                                          cytostatic;
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                                                                      Zlotnik A;
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                                                                                           INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for treating leukaemia.
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Early detection of soft tissue sarcoma comprises determining expression of a gene in a first soft tissue sample and a normal soft tissue sample and comparing the gene expression, also useful in treating soft tissue

tissue sample soft tissue

sarcoma.

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Matches
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Best Local
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which comprises obtaining a first soft tissue sample from an individual and a normal soft tissue sample from the same or different individual, determining the expression of a gene in both samples and comparing the expression of the gene in both soft tissue samples, where a higher level of protein expression in the first soft tissue sample indicates the presence of soft tissue sarcoma. The method of the invention has cytostatic applications and may be useful for detecting soft tissue sarcoma, possibly via gene therapy or vaccine production. The nucleic
                                                                                                                          The invention relates to a novel method for detecting soft tissue sarcoma which comprises obtaining a first soft tissue sample from an individual
                                                                                                                                                                                                                                     Early detection of soft tissue sarcoma comprises determining expression of a gene in a first soft tissue sample and a normal soft tissue sample and comparing the gene expression, also useful in treating soft tissue
                                                                                                                                                                                                                                                                                                                                                 Aziz N,
                                                                                                                                                                                                                                   of a gene in a tra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      soft tissue sarcoma;
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                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-NOV-2003; 2003WO-US038193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004048938-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1797 BP; 574 A; 340 C; 360 G; 523 T;
                                                                                                                                                                                                                                                                                                                                                                                    (PROT-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention relates to a novel method for detecting soft tissue sarcoma ch comprises obtaining a first soft tissue sample from an individual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                  PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                               Ginsburg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGACCCTAGAGGCAAAG
                                                                                                                                                                                  SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               85.6%;
94.1%;
                                                                                                                                                                                896; 210pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic; gene therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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Pred. No. 5.
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                                                                                                                                                                                  English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    screening;
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RESULT 22
ADZ59509/G
ID ADZ595
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                                                               involves detecting a variation in a gene chosen from CACNAIC, CALCRL, CHI3L1, EGF, FGF1, GFRA1, GPR56, GPRK6, IL10RA, IL10RB, IL12RB1, KCNJ14, KCNJ1, ORCTL4, PDGFRA, SCYB14, SLC12A1, SLC12A3, TGFBR3, TMEM1, CALCR, IL17R, OSTP1, FGF6, HGF, MET, TGFB1 and VEGF, or detecting the base in a polymorphism region existing in the vicinity of any one of the genes. The invention further comprises a reagent or kit for testing secondary hyperparathyroidism in a chronic renal failure patient. This polymorphism cequence represents the polymorphism containing human CHI3L1 gene of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Testing secondary hyperparathyroidism in chronic involves detecting variation in gene chosen from EGF, FGF1, GFRAL, GFR56 and GFRK6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid sequences may be useful in diagnostic and screening applications. The current sequence is that of a human soft tissue sarcoma-upregulated DNA of the invention. The current sequence is not shown within the specification per se but was submitted in CD format by the inventor.
                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel method for testing secondary hyperparathyroidism in a chronic renal failure patient. The method
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 3; 138pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-SEP-2003; 2003JP-00341015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-SEP-2003; 2003JP-00341015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP2005102601-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        secondary hyperparathyroidism; endocrine-gen.; antithyroid;
renal failure; nephrotropic; SNP detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Secondary hyperparathyroidism detection human polymorphic gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JUN-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HYUB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      319
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HYUBITTO GENOMICS KK UNIV JIKEI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9645 BP; 2115 A; 2895 C; 2715 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nephrotropic; SNP detection de polymorphism; SNP; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /standard_name= "Single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             85.6%;
94.1%;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            renal failure patient, CACNA1C, CALCRL, CHI31
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Query Match
Best Local Similarity
Matches 16; Conserv

Conservative

85.6%; 94.1%;

Pred. No. 5.8e+02; 0; Mismatches 1

Indels

0,

Gaps

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Score 15.4; Pred. No. 5.

DВ

Sequence 20001 BP; 4773 A; 5041 C;

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5029

T; 0 U; 0 Other;

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RESULT 23
ADQ97998/c
ID ADQ97998 standard; DNA; 22
XX
AC ADQ97998;
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AC ADQ97998;
XX
DT 07-OCT-2004 (first entry)
XX
DE Human cancer associated se
XX
Cytostatic; Gene Therapy;
XX
DE Human cancer associated se
XX
Cytostatic; Gene Therapy;
XX
DE 2-JUL-2004.
XX
PM W02004060304-A2.
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PM 22-JUL-2004.
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PM 22-DEC-2003; 2003WO-US041:
XX
PF 22-DEC-2002; 2002US-00330'
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PF 27-DEC-2002; 2002US-00330'
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PF 27-DEC-2002; 2002US-00330'
XX
PM (SAGR-) SAGRES DISCOVERY:
XX
PM (SAGRES D
                                                                                                                                                                                                      RESULT 24
ADM69029
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to cancer associated sequences (ADQ97025-ADQ98004). The sequences are useful for the diagnosis, prevention and/or treatment of cancer, such as leukemia and lymphoma. Note: The sequence data for this patent did not form part of the printed specification, but we shall did not formate directly from WIPO at
                                          platelet derived growth factor receptor alpha; PDGFRA; neoplasia; human;
wild-type; type III receptor tyrosine kinase; RTK; ds; gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated cancer associated nucleic acids comprising at least 10 contiguous nucleotides, useful for diagnosing, preventing and/or treating cancers such as leukemia and lymphoma.
                                                                                           Human platelet derived growth factor receptor alpha wild-type gDNA
                                                                                                                           03-JUN-2004
                                                                                                                                                          ADM69029;
                                                                                                                                                                                       ADM69029 standard; DNA; 191150 BP
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 22475 BP; 5898 A; 5414 C; 5434 G;
                                                                                                                                                                                                                                                                                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 975; 199pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cytostatic; Gene Therapy; cancer; leukemia; lymphoma; Human; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human cancer associated sequence HD11-038,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SAGR-) SAGRES DISCOVERY INC
                                                                                                                                                                                                                                                                 15390 GGACCCTAGAGGAAAAG 15374
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nilarity 94.1%;
Conservative
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Pred. No. 5.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                              5709 T; 0 U; 20 Other;
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lpha wild-type protein"
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RESULT 25
ABZ07598
ID ABZ077
XX ABZ077
XX O9-JA
XX T7; 1
KW T7; 1
KW T7; 1
KW Theum
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Best Local (
                                                                                  New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               T7; leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;
                                                                                                                                                   WPI; 2002-636525/68.
                                                                                                                                                                                                   Wohlgemuth J,
                                                                                                                                                                                                                                                                 20-OCT-2000; 2000US-0241994P
08-JUN-2001; 2001US-0296764P
                                                                                                                                                                                                                                                                                                                   22-OCT-2001; 2001WO-US047856
                                                                                                                                                                                                                                                                                                                                                   25-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                  WO200257414-A2
                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human leukocyte gene expression profiling probe SEQ ID NO 7589.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ07598;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 191150 BP; 55223 A; 39036 C; 39352 G; 56412 T; 0 U; 1127 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genomic DNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                factor receptor alpha (PDGFRA) polypeptide. The polypeptide of the invention is a type III receptor tyrosine kinase (RTK) and may be useful for screening for a compound that may influence PDGFRA-mediated neoplasia in a mammal. The current sequence is that of the human PDGFRA wild-type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated variant platelet derived growth factor alpha (PDGFRA) polypeptide, useful for screening for a compound useful in influencing PDGFRA-mediated neoplasia in a mammal.
                                                                                                                                                                                                                                  (BIOC-) BIOCARDIA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel isolated variant platelet derived growth
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P-PSDB; ADM69012.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   117726
                                                                                                                                                                                muth J, Fry K,
Woodward R, Qu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA; 50
                                                                                                                                                                                  K, Matcuk G, I
Quertermous T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  85.6%;
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Pred. No. 6.2e+02.
0; Mismatches
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The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of

Claim 1;

Page 572; Opp; English.

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RESULT 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50-mer oligonucleotide marker probe of the invention #229
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20-DEC-2002;
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2002US-00325899.
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Pred. No.
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7.6e+02;
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The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic slet, lung, bone marrow or stem cell transplant rejection, in an individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lung, bacterial or fungal infection. The present sequence represents a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft

an

Query Match Best Local Similarity

83.3%; 100.0%;

Score 15; Pred. No. G; 11 T;

51;

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P

12 ü 19

0 U; 0 Other; DB 4; Length 7.6e+02;

Claim 2;

SEQ ID NO 229; 1762pp; English.

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cu polymucleotide sequences. The sequences can be used in gene and protein contempy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases content and be used in the prevention, diagnosis and treatment of diseases content of the prevention, diagnosis and treatment of diseases content of the prevention of polypeptides. For content of the prevention of polypeptides or content of the activity of polypeptides content of the production of polypeptides. Additionally, (I) and its complementary contents own content of polypeptides and the presence of similar nucleic acids in samples, and content the prevention of polypeptides of similar nucleic acids in samples, and content of polypeptides encoded by (I) may be used as antigens in the production of polypeptides encoded by (I) may be used as antigens in the production of the used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic contents of polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic contents of polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic contents may be used as diagnostic agents for detecting the presence of polymorphic contents may also be used as diagnostic agents for detecting the presence of polymorphic contents may be used as diagnostic agents for detecting the presence of polymorphic contents may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic contents may be used as diagnostic agents for detecting the presence of polymorphic contents may be used as diagnostic agents for detecting the presence of polymorphic contents may be used as diagnostic agents for detecting the presence of polymorphic contents may be used as diagnostic agents for detecting the presence of polymorphic contents may be used as diagn
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AAI76987/c
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                                                                                                                                                                                                                                                                                                                                                                                               AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein polynucleotide sequences.
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 1254; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-356160/37
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100.0%; Pr
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k; Pred. No. 7.6
0; Mismatches
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detection;
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                                                         Query Match
Best Local S
Matches 15
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                                                                                                                                                                                                              by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, [1] and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by [1] may be used as antigens in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6986/c
                                                                                                                                                                  antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                        therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                       Sequence 51
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                     polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                       sequences (I), which
AAM53114 to AAM53329
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 1253; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphic nucleic acid sequences, useful in genetic testing
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                            2 GGACCCTAGAGGCAA 16
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                                                                          Similarity
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                                                                                                                       B₽,
                                                         83.3%; Sillarity 100.0%; Conservative 0;
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                                                                                                                       A;
                                                                                                                       12 C; 20
                                                                                                                                                                                                                                                                                                                                                                                                                    contain single nucleotide polymorphisms
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                                                                          Score 15;
Pred. No.
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                                                             Mismatches
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                                                      DB 4; L-.
3. 7.6e+02;
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                                                                                        Length 51;
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detection;
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RESULT 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to novel human nucleic acid sequences (I). CC The present sequence is one such nucleic acid sequence. Expression of (I) CC are implicated in tumour suppression or reversion and apoptosis and viral CC resistance. (I) are useful as probes or primers for detecting, CC identifying, measuring and/or amplifying nucleic acid sequences, as CC entisense reagents and for recombinant production of polypeptides. (I), CC polypeptides (II) encoded by (I), vector containing (I), cells containing CC these vectors and antibodies (Ab) against (II) are all useful for CC (especially neurodegeneration, such as Alzheimer's disease and CC (especially neurodegeneration). Nuch as Alzheimer's diseases (also immune CC (i) are used for studying the actiology of these diseases (also immune CC and inflammatory diseases). Note: In the present specification, SCQ ID 1 to 2280 are claimed in Claim 1, however only SEQ ID 1 to 2270 are shown or the specification.
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 75 BP; 27 A; 12 C;
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06-OCT-2000 AAC24825;

(first entry)

Human secreted protein 5'

EST, SEQ ID NO: 28900.

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                                                                                                                                                               Query Match 83.3%; Score 15; DB 3; Lu
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                          The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed CDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer CDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
                                                                                                                                                                                                                                                    Sequence 163 BP; 35 A; 57 C; 41 G; 30 T; 0 U; 0 Other;
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Sequence 55145, Application US/10301480 Publication No. US20060057564A1 GENERAL INFORMATION:
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RESULT 1 US-10-301-480-55145

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APPLICANY: Wang, David G.
TITLE OF INVENTION: Identifiction and Mappin
TITLE OF INVENTION: in the Human Genome
FILE REFERENCE: 108827.137
CURRENT APPLICATION NUMBER: US/10/301,480
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US 10/215,598
PRIOR FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: US 60/311,695
PRIOR APPLICATION NUMBER: US 60/311,695
PRIOR FILING DATE: 2001-08-10
NUMBER OF SEQ ID NOS: 1226818
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 668554
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TITLE OF INVENTION: Identifiction and Mapping of
TITLE OF INVENTION: in the Human Genome
FILE REFERENCE: 108827.137
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/10/301,480
CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: US 10/215,598
PRIOR PILING DATE: 2002-08-09
PRIOR PILING DATE: 2002-08-09
PRIOR FILING DATE: 2001-08-10
NUMBER OF SEQ ID NOS: 1226818
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 55145
LENGTH: 539
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; Sequence 504327, Application US/09925065A
; Publication No. US20040181048A1
; GENERAL INFORMATION:
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US-10-301-480-668554
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US-10-301-480-668554
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; ORGANISM: Homo sapien
US-10-301-480-55145
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APPLICANT: Wang, David G.
TITLE OF INVENTION: Nucleotide Polymorphisms in the Human
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR APPLICATION NUMBER: US 60/243,096
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Pred. No. 2.7e
0; Mismatches
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Genome
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-925-065A-504327
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US-09-925-065A-65294
Sequence 65295, Application US/09925065A
Publication No. US20040181048A1
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Nucleotide Polymorphisms in the Human Genome
FILE REFERENCE: 108827.135
CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
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APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Nucleotide Polymorphisms in the Huma
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PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: US 60/289,846
PRIOR FILING DATE: 2001-05-09
NUMBER OF SEQ ID NOS: 957086
SOFTMARE: FASTSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR FILING DATE: 2000-11-30
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR FILING DATE: 2001-01-16
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PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR FILING DATE: 2000-10-24
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CURRENT APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
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PRIOR APPLICATION NUMBER: US 60/250,092
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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Local Similarity 94.1%;
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94.1%;
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Pred. No. 2
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Pred. No. 2.7e+02;
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; ORGANISM: Homo sapiens
US-09-925-065A-65295
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US-10-301-480-166533/c
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GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Ident
TITLE OF INVENTION: in t
                                                                        Sequence 166533, Application US/10301480 Publication No. US20060057564A1
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LENGTH: 738
TYPE: DNA
ORGANISM: Homo sapiens
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TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Nucleotide Polymorphisms in the Huma
FILE REFERENCE: 108827.135
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SOFTWARE: FastSEQ for Windows Version 4.0
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OR FILING DATE: 2000-10-24
OR APPLICATION NUMBER: US 60/252,147
OR APPLICATION NUMBER: US 60/250,092
OR APPLICATION NUMBER: US 60/250,092
OR FILING DATE: 2000-11-30
OR APPLICATION NUMBER: US 60/261,766
OR FILING DATE: 2001-01-16
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94.1%;
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0; Mismatches
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Pred. No. 2.
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Identifiction and Mapping

of Single Nucleotide Polymorphisms

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RESULT 9
US-10-301-480-166535/c
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US-10-301-480-166533
                                                                                                                                                                     Sequence 166535, Application US/10301480
Publication No. US20060057564A1
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identifiction and Mapping of Single Nucleotide Polymorphisms
TITLE OF INVENTION: in the Human Genome
FILE REFERENCE: 108827.137
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 166534
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-10-301-480-166534
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                                                                              CURRENT APPLICATION NUMBER: US/10/301,480
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US 10/215,598
PRIOR FILING DATE: 2002-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2002-11-21 PRIOR APPLICATION NUMBER: US 10/215,598 PRIOR FILING DATE: 2002-08-09
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TITLE OF INVENTION: Identifiction and Mapping of Single Nucleotide Polymorphisms
TITLE OF INVENTION: in the Human Genome
                    PRIOR APPLICATION NUMBER: US 60/311,695
PRIOR FILING DATE: 2001-08-10
NUMBER OF SEQ ID NOS: 1226818
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PRIOR APPLICATION NUMBER: US 10/215,598
PRIOR FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: US 60/311,695
PRIOR FILING DATE: 2001-08-10
PRIOR FILING DATE: 2001-08-10
NUMBER OF SEQ ID NOS: 1226818
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CURRENT FILING DATE: 2002-11-21
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FastSEQ for Windows Version 4.0
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Pred. No. 2.8e+02
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Pred. No. 2.8e+02;
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; ORGANISM: Homo sapien 
US-10-301-480-779943
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; ORGANISM: Homo sapien
US-10-301-480-166535
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                                                                                                                           SOFTWARE: FastSEQ for Windows
SEQ ID NO 779943
LENGTH: 738
                                                                                                                                                                                                                                                                                                                                  Sequence 779943, Application US/10301480
Publication No. US20060057564A1
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identifiction and Mapping of Single Nucleotide Polymorphisms
TITLE OF INVENTION: in the Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 1226818
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 779942
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Best Local (
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                                   Query Match
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PRIOR FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: US 60/311,695
PRIOR FILING DATE: 2001-08-10
                                                                                                                                                                                                                                                                             FILE REFERENCE: 108827.137
CURRENT APPLICATION NUMBER: US/10/301,480
CURRENT FILING DATE: 2002-11-21
                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
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CURRENT FILING DATE: 2002-11-21
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TITLE OF INVENTION: Identifiction and Mapping of Single Nucleotide Polymorphisms
TITLE OF INVENTION: in the Human Genome
FILE REFERENCE: 108827.137
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                 Local Similarity
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94.1%;
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94.1%;
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Pred. No. 2
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Score 15:4; DB 10
Pred. No. 2.8e+02;
0; Mismatches 1
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Pred. No. 2
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Matches

16;

Conservative

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Gaps

GGACCCTAGAGGCAAAG

18

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FILE REFERENCE: MMI1100-2
CURRENT APPLICATION NUMBER: US/10/750,185
CURRENT FILING DATE: 2003-12-31
PRIOR APPLICATION NUMBER: US 60/437,482
PRIOR FILING DATE: 2002-12-31
NUMBER OF SEQ ID NOS: 64922
SOPTWARE: PatentIN version 3.1
SEQ ID NO 61166
LENGTH: 1155
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US-10-750-185-61166
; Sequence 61166, Application US/10750185
; Publication No. US20050260603A1
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US-10-301-480-779944
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SOFTWARE: FRETSEQ for Windows Version 4.0
SEQ ID NO 779944
LENGTH: 738
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                                                                              Matches
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TITLE OF INVENTION: Identifiction and Mapping of Single Nucleotide Polymorphisms
TITLE OF INVENTION: in the Human Genome
FILE REFERENCE: 108827.137
CURRENT APPLICATION NUMBER: US/10/301,480
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US 10/215,598
PRIOR APPLICATION NUMBER: US 10/215,598
PRIOR APPLICATION NUMBER: US 60/311,695
PRIOR APPLICATION NUMBER: US 60/311,695
PRIOR FILING DATE: 2001-08-10
PRIOR FILING DATE: 2001-08-10
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Publication No. US20060057564A1
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TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MMI GENOMICS, INC.
                                                                                                                                                                                                 TYPE: DNA
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                                                                            Local Similarity es 16; Conserv
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GGACCCTAGAGACAAAG 885
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HOLM, Tom
BATES, Stephen
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                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Richard
                                                                                                                                                                            19866881037774
                                                                                              85.6%; Score 15.4;
94.1%; Pred. No. 2.
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Pred. No. 2
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                                                                                                  2.9e+02
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RESULT 16
US-10-932-182A-2596/c
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SEQ ID NO 2596
SERVID: 1206
TYPE: DNA
ORGANISM: Saccharomyces pastorianus
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Publication No. US20060046253A1
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: FUJIMURA, TOMOKO
APPLICANT: ASHLKARI, TOSHIHIKO
APPLICANT: ASHLKARI, TOSHIHIKO
TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS
FILE REFERENCE: 030685-043
FULL REFERENCE: 030685-043
CURRENT APPLICATION NUMBER: US/10/932,182A
CURRENT FILLING DATE: 2004-09-02
NUMBER OF SEQ ID NOS:.197023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/437,482
PRIOR FILING DATE: 2002-12-31
NUMBER OF SEQ ID NOS: 64922
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CURRENT APPLICATION NUMBER: US/10/750,623
CURRENT FILING DATE: 2003-12-31
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APPLICANT: ROSENFELD, David
APPLICANT: HOLM, Tom
APPLICANT: BATES, Stephen
APPLICANT: FANTIN, Dennis
TITLE OF INVENTION: METHODS AND SYSTEMS FOR INFERRING BOVINE TRAITS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: NAKAO,
APPLICANT: NAKAM
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TYPE: DNA
ORGANISM: Bovine 19866881037774
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APPLICANT: MMI GENOMICS, INC.
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Local Similarity 94.1%;
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KODAMA, YUKIKO
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94.1%;
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Pred. No. 2.9e+02;
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Sequence 2596, Application US/10932182A publication No. US20060046253A1

GENERAL INFORMATION

NAKAMURA, NORIHISA

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                                                                                            Sequence 2408, Application US/11096568A

Publication No. US20060048240A1

GENERAL INFORMATION:

APPLICANT: Alexandrov, Nickolai et al.

TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides

TITLE OF INVENTION: Therby

FILE REFERENCE: 2750-1592PUS2

CURRENT APPLICATION NUMBER: US/11/096,568A

CURRENT APPLICATION NUMBER: US/11/096,568A

CURRENT FILING DATE: 2005-04-01

NUMBER OF SEQ ID NOS: 34471

SEQ ID NO 2408
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LENGTH: 1206
TYPE: DNA
ORGANISM: Saccharomyces pastorianus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides TITLE OF INVENTION: Therby FILE REFERENCE: 2750-1592PUS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS FILE REFERENCE: 030685-043
CURRENT APPLICATION NUMBER: US/10/932,182A
CURRENT FILING DATE: 2004-09-02
NUMBER OF SEQ ID NOS: 197023
SOFTWARE: Patentin version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 34471
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(1806)
OTHER INFORMATION: Ceres Seq. ID no. 13603467
                                       TYPE: DNA
ORGANISM: Arabidopsis thaliana
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AME/KEY: misc_feature
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NUMBER OF SEQ ID NOS: 7285
SOFTWARE: PatentIn version 3.3
SEQ ID NO 856
LENGTH: 8703
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Publication No. US20060003958A1
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CURRENT FILING DATE: 2005-05-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sinacore, Mark W.
APPLICANT: Leonard, Mark W.
APPLICANT: Brown, Eugene L.
APPLICANT: Miller, Christopher P.
APPLICANT: Miller, Christopher P.
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES RELATED TO OLIGONUCLEOTIDE ARRAYS
TITLE OF INVENTION: TO MONITOR GENE EXPRESSION
FILE REFERENCE: 01997.027701
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OTHER INFORMATION: Ceres Seq. ID no. 14312112
NAME/KEY: misc feature LOCATION: (2778)..(2795) OTHER INFORMATION: n is
                                                              NAME/KEY: misc feature
LOCATION: (1728)..(1776)
OTHER INFORMATION: n is
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LOCATION: (1384)..(145
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LOCATION: (331)..(349)
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ORGANISM: Mesocricetus auratus
                                                                                                                           LOCATION: (1384)..(1455)
OTHER INFORMATION: n is
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OTHER INFORMATION: n is
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NAME/KEY: misc_feature
LOCATION: (402)..(414)
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OTHER INFORMATION: n is
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LOCATION: (417)..(446)
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WAME/KEY: misc\_feature

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APPLICANT: Sinacore, Martin S.
APPLICANT: Leonard, Mark W.
APPLICANT: Leonard, Mark W.
APPLICANT: Miller, Christopher P.
APPLICANT: Miller, Christopher P.
TITLE OF INVENTION: OLICONUCLEOTIDE ARRAYS TO
TITLE OF INVENTION: MAKING AND USING SAME
FILE REFERENCE: 01997.027700
CURRENT APPLICATION NUMBER: US/11/128,049
CURRENT FILING DATE: 2005-05-11
PRIOR APPLICATION NUMBER: US 60/570,425
PRIOR FILING DATE: 2004-05-11
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JS-11-128-049-856/c
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Best Local Similarity
Matches 16; Conserv
SOFTWARE: PatentIn
SEQ ID NO 856
LENGTH: 8703
                                                                                                                                                                                                                                                                                                       Sequence 856, Application No.
                                          NUMBER OF SEQ ID NOS:
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APPLICANT: Charlebois, Timo
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LOCATION: (8381)..(841
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LOCATION: (7932)..(7949)
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THER INFORMATION: n is
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LOCATION: (4437)..(4454)
OTHER INFORMATION: n is a,
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OTHER INFORMATION: n is a,
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OTHER INFORMATION: n is a,
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JOCATION: (7677)..(7701)
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OTHER INFORMATION: n is a,
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OCATION: (4895)..(492)
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OCATION: (3287)..(334
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THER INFORMATION: n is a,
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o. US20060010513A1
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PEATURE:

NAME/KEY: misc feature

TOTATION: (331)...(349)
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LOCATION: (402)..(414)
OTHER INFORMATION: n is
FEATURE:
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NAME/KEY: misc_feature
LOCATION: (279)...(291)
OTHER_INFORMATION: n is a,
                               FEATURE: misc feature
LOCATION: (7932)..(7949)
OTHER INFORMATION: n is
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LOCATION: (7861)..(7903)
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NAME/KEY: misc feature

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OTHER INFORMATION: n is
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LOCATION: (4895)..(4922)
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LOCATION: (3342)...(3362)
OTHER INFORMATION: n is
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NAME/KEY: misc_feature
LOCATION: (2778)...(2795)
OTHER INFORMATION: n is
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OTHER INFORMATION: n is
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ORGANISM: Mesocricetus
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OTHER INFORMATION: n is
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LOCATION: (4437).:(445
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NAME/KEY: misc_feature
NOCATION: (8381)..(8418)
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OCATION: (3287)...(334
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-124-367A-5024
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US-11-124-367A-5024
                                                                                                                                                             CURRENT APPLICATION NUMBER: US/11/124,367A
CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US 60/568,846
PRIOR FILING DATE: 2004-05-07
PRIOR APPLICATION NUMBER: US 60/582,609
PRIOR PILING DATE: 2004-06-25
PRIOR APPLICATION NUMBER: US 60/599,554
PRIOR PILING DATE: 2004-08-09
NUMBER OF SEQ ID NOS: 34460
SOFTWARE: FBSLSEQ for Windows Version 4.0
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Matches
                                                                                                                            SEQ ID NO 5024
LENGTH: 100000
Query Match 85.6%;
Best Local Similarity 94.1%;
Matches 16; Conservative
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Best Local Similarity 94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                           APPLICANT: Michele Cargill
APPLICANT: Hongin Huang
TITLE OF INVENTION: Fehretic Polymorphisms Associated with
TITLE OF INVENTION: Fibrosis Methods of Detection and Uses Thereof
FILE REFERENCE: CL001519.ORD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/330,773
CURRENT FILING DATE: 2002-12-27
NUMBER OF SEQ ID NOS: 981
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Marc Malandro
TITLE OF INVENTION: Novel Compositions and Methods in Cancer
FILE REFERENCE: 529452001300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
PAME/KEY: misc_feature
LOCATION: (1)...(22475)
OTHER INFORMATION: n = A,T,C or G
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94.1%;
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Pred. No. 3.7e+02;
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Pred. No. 3.4e+02
                 Score 15.4; DB 14
Pred. No. 4.1e+02;
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 RESULT 25
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US-11-101-244-1292346
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; Sequence 1292346, Application US/11101244

; Publication No. US20050246794A1
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Best Local Similarity
Watches 16; Conserve
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SEQ ID NO 1292346
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                                                                                                                                             Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/502,050
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/426,137
PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 1591911
                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/11/101,244
CURRENT FILING DATE: 2005-04-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/11/121,086
CURRENT FILLING DATE: 2005-05-04
PRIOR APPLICATION NUMBER: 60/567,570
PRIOR FILLING DATE: 2004-05-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Marshall, William
APPLICANT: Scaringe, Stephen
TITLE OF INVENTION: Functional and Hyperfunctional sirNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dharmacon, APPLICANT: Khvorova,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: POULSEN, TIM S.

APPLICANT: NIELSEN, KIRSTEN V.

TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES FILE REFERENCE: 09138.6000-00000
                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 13499US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 107
                                                                                                                                                                                                                                         TYPE: RNA
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                           LENGTH: 19
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                                                       GGACCCUAGAGGCAA 18
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Reynolds, Angela
                                                                                                                              Conservative
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94.1%;
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Pred. No.
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Pred. No. 4.3e+02;
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DB 14;

Length 171732;

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DB 12; .3e+02;

Length 19; Indels

<u>.</u>.

Gaps

US-11-101-244-1292367

ce 1292367, Application US/11101244 ation No. US20050246794A1

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US-11-083-784-1292346
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US-11-083-784-1292346
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US-11-101-244-1292367
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PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 1591911
SOFTWARE: Proprietary
SEQ ID NO 1292367
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                                    Matches
                                                                       Query Match
                                                                                                                                                                                     SEQ ID NO 1292346
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TITLE OF INVENTION: Functional and Hyperfunctional sirNA
FILE REFERENCE: 13499US
CURRENT APPLICATION NUMBER: US/11/083,784
CURRENT FILING DATE: 2005-03-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/426,137 PRIOR FILING DATE: 2002-11-14
                                                                                                                                                                                                                                                                          CURRENT FILING DATE: 2005-03-18
PRIOR APPLICATION NUMBER: US/10/714,333
PRIOR FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR FILING DATE: 2003-09-10
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CURRENT ETLING DATE: 2005-04-07
CRICK APPLICATION NUMBER: 60/502,050
PRIOR FILING DATE: 2003-09-10
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TITLE OF INVENTION: Functional and Hyperfunctional siRNA
FILE REFERENCE: 13499US
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APPLICANT: Khvorova,
APPLICANT: Reynolds,
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2 GGACCCTAGAGGCAA 16
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Similarity 93.3%;
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Reynolds, Angela
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Marshall, William
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Reynolds, Angela
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                                    Conservative
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93.3%;
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Pred. No. 3
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Pred. No. 3.3e+02;
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                                                                       DB 13; Length 19;
                                                       .3e+02;
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US-11-083-784-1292367
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                                                              RESULT 29
                                                                                                                                                                                                                                                                         ; ORGANISM: Human
US-10-310-914A-965984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-310-914A-965984/c
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 965984
LENGTH: 22
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Sequence 799050, Application US/09925065A Publication No. US20040181048A1
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SEQ ID NO 1292367
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Publication No. US20050245475A1
                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Bioinformatically detectable group of novel regulatory genes and TITLE OF INVENTION: uses thereof FILE REFERENCE: 06087.0200.CPUS01 CURRENT APPLICATION NUMBER: US/10/310,914A CURRENT FILING DATE: 2002-12-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/11/083,784
CURRENT FILING DATE: 2005-03-18
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PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 1591911
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PRIOR FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/502,050
PRIOR FILING DATE: 2003-09-10
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APPLICANT: Scaringe, Stephen
IITLE OF INVENTION; Functional and Hyperfunctional siRNA
                                                                                                                                                                                                                                                                                                               TYPE: RNA
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Reynolds, Angela
Leake, Devin
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APPLICANT: Wang, David G.

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Search completed: April 15, 2006, 18:19:53
Job time : 425 secs
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 799051
LENGTH: 640
TYPE: DNA
CRGANISM: Homo sapiens
US-09-925-065A-799051
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; ORGANISM: Homo sapiens
US-09-925-065A-799050
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PRIOR APPLICATION NUMBER: US/09/925,065A
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR APPLICATION NUMBER: US 60/252,147
PRIOR FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR FILING DATE: 2001-01-16
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CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: US 60/243,096
PRIOR FILING DATE: 2000-10-24
PRIOR PAPLICATION NUMBER: US 60/252,147
PRIOR FILING DATE: 2000-11-26
PRIOR APPLICATION NUMBER: US 60/250,092
PRIOR FILING DATE: 2000-11-30
PRIOR FILING DATE: 2000-11-30
PRIOR APPLICATION NUMBER: US 60/261,766
PRIOR PILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: US 60/269,846
PRIOR PILING DATE: 2001-05-09
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Publication No. US20040181048A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2001-05-09
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TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Nucleotide Polymorphisms in the Human Genome
FILE REFERENCE: 108827.135
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41 14 //.8 201 8 US-10-719-993	40 14 77.8 201 7 US-10-741-6	38	37 14.4 80.0 3186778 6 US-10-027-	35	234 14.4 80.0 169567 8 US-10-719-9	232 14.4 80.0 142976 7 US-10-367-0	231 14.4 80.0 127767 7 US-10-322-2	30 14.4 80.0 116704 5 US-10-087-1	29 14.4 80.0 115218 8 US-10-278-6	228 14.4 80.0 115218 8 US-10-278-6	226 14.4 80.0 103747 7 US-10-450-8	225 14.4 60.0 11950 / US-10-087-1	24 14.4 80.0 18443 3 US-US-741-6	223 14.4 80.0 8323 3 US-09-970-5	222 14.4 80.0 7036 5 US-10-178-7	221 14.4 80.0 3153 6 US-10-104-0	220 14.4 80.0 2754 3 US-09-938-8	219 14.4 80.0 2754 3 US-09-938-8	18 14.4 80.0 2621 10 US-11-097-	15 14.4 80.0 2136 / US-10-424-51	215 14.4 80.0 1354 4 05-09-9495-10	214 14.4 80.0 1334 4 US-09-925-04	213	213 14 4 80 0 1262 4 00 0 27 00 0	212 14.4 80.0 1262 4 US-09-925-00	11 14 4 80.0 939 8 US-10-926-68	210 14 4 80 0 939 6 US-10-170-0	200 14.4 80.0 878 6 US-10-027-6	207 14.4 80.0 878 5 US-10-027-6	05 14.4 80.0 040 0 03-10-027-5	05 14.4 80.0 845 6 IIS-10-027-6	05 14.4 80.0 845 5 US-10-027-6	0.0 14.4 00.0 010 1 00 00 000 00	02 14.4 00.0 013 4 00-09-925-06	OF 14.4 00.0 013 4 00-03-363-04	00 14.4 00.0 733 / 00.40 424 0.	200 14 4 80 0 733 7 IIS-10-424-50	199 14.4 BO 0 703 6 US-10-027-6	198 14.4 80.0 703 6 US-10-027-63	197 14.4 80.0 703 6 US-10-027-6	196 14.4 80.0 703 6 US-10-027-6	195 14.4 80.0 703 5 US-10-027-60	194 14.4 80.0 703 5 US-10-027-63	193 14.4 80.0 703 5 US-10-027-63	192 14.4 80.0 703 5 US-10-027-63	14.4 80.0 663 6 US-10-027-6	14.4 80.0 663 5 US-10-02/-61	14.4 80.0 658 5 173 673 675	14.4 80.0 658 6 05-10-1/0-03	14.4 80.0 636 8 08-10-33/13/1	186 14.4 80.0 632 6 US-10-02/-61	185 14.4 80.0 632 6 US-10-02/-61	184 14.4 80.0 632 5 US-10-02/-6	183 14.4 80.0 632 5 US-10-02/-61	182 14.4 80.0 621 10 057.1.	161 14.4 60.0 610 0 05-t0-02/-02	10 11 1 00 0 010 0 00 10 017 01	180 14 4 .80 0 .616 6 IIS-10-027-6	179 14.4 80.0 616 5 US-10-027-61	178 14.4 80.0 616 5 US-10-027-63	T4.4 80.0 SEC 5 CS-10-02/-01	14 4 00 0 610 6 TE-10-027-63	76 14.4 80.0 610 6 US-10-027-63	75 14.4 80.0 610 6 US-10-027-6:	74 14.4 80.0 610 5 US-10-027-6	14.4 80.0 610 5 US-10-027-6:	14.4 80.0 610 5 US-10-027-6	171 14.4 80.0 610 4 US-09-925-08	170 14.4 80.0 606 4 US-09-925-06	120 11 1 200 COC 1 TTC-09-925-06
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41 14 77.8 201 8 US-10-719-993-22356	14 77.8 201 7 US-10-741-601-7861	38 14 77.8 25 8 US-10-719-900-763654	37 14.4 80.0 3186778 6 US-10-027-632-174961	35 14.4 80.0 329019 8 US-10-388-838-48	234 14.4 80.0 169567 8 US-10-719-993-6774	232 14.4 80.0 142201 8 05-10-367-094-99 233 14.4 80.0 142976 7 US-10-367-094-99	231 14.4 80.0 127767 7 US-10-322-281-797	30 14.4 80.0 116704 5 US-10-087-192-1456	29 14.4 80.0:115218 8 US-10-278-698-769	228 14.4 80.0 115218 8 US-10-278-698-255	220 14.4 80.0 103747 7 US-10-450-826-46	225	24	223 14.4 80.0 8323 3 US-09-970-597-2	222 14.4 80.0 7036 5 US-10-178-782-3	221 14.4 80.0 3153 6 US-10-104-047-327	220 14.4 80.0 2754 3 US-09-938-842A-1786	219 14.4 80.0 2754 3 US-09-938-842A-1786	18 14.4 80.0 2621 10 US-11-097-143-36034	15	215	214	0.0 10.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0	212 14.4 80.0 1262 4 US-09-925-065A-727392 9	212 14.4 80.0 1262 4 US-09-925-065A-727391	11 14 4 80.0 939 8 US-10-926-684-403	210 14 4 80 0 939 6 US-10-170-097-403	209 14.4 80.0 878 6 US-10-027-632-136896	20/ 14.4 00.0 040 0 05-10-027-032-136896 0	05 14.4 00.0 040 0 00-10-07-30-157313 0	05 14.4 80.0 845 6 IIS=10=027=632=157312 8	04 14.4 80.0 845 5 US-10-027-632-157313 C	03 14.4 80.0 845 5 IIS-10-027-632-157312 8	02 14.4 00.0 e13 / 05-09-921-0653-7232-7	OL 14.4 GO.O GIJ 4 CO-OG-26-26-73337 (	00 14.4 00.0 733 7 00.40 224 077 110071	200 14 4 80 0 733 7 15-10-424-599-136542	199 14.4 RO.0 703 6 US-10-027-632-25471	198 14.4 80.0 703 6 US-10-027-632-25470	197 14.4 80.0 703 6 US-10-027-632-25469	196 14.4 80.0 703 6 US-10-027-632-25468	195 14.4 80.0 703 5 US-10-027-632-25471	194 14.4 80.0 703 5 US-10-027-632-25470	193 14.4 80.0 703 5 US-10-027-632-25469	192 14.4 80.0 703 5 US-10-027-632-25468	14.4 80.0 663 6 US-10-027-632-227862	14.4 80.0 663 5 US-10-02/-632-22/862	14.4 80.0 658 8 US-10-926-684-402	14.4 80.0 658 6 05-10-1/0-09/-402	14.4 80.0 636 8 08-10-30/-300-00460	14.4 80.0 632 6 05-10-02-20-20-20-20-20-20-20-20-20-20-20-20	185 14.4 80.0 632 6 US-10-02/-632-2/5520	184 14.4 80.0 632 5 US-10-02/-632-2/5921	183 14.4 80.0 63.5 10-10-02/-632-2/3320	182 14.4 80.0 621 10 05-11-09/-140-0000	100 14.4 00.0 00 0 00 10 10 10 10 10 10 10 10 10 10	10 14 1 00 0 616 6 IIs-10-027-632-307702	180 14 4 .80 0 616 6 IIS-10-027-632-93222	179 14.4 80.0 616 5 US-10-027-632-307702	: 178 14.4 80.0 616 5 US-10-027-632-93222 S	14.4 00.0 otto 00"10"00"0	14 4 60 0 630 6 IIS-10-037-633-234707 S	.76 14.4 80.0 610 6 US-10-027-632-224706 S	75 14.4 80.0 610 6 US-10-027-632-224705 S	74 14.4 80.0 610 5 US-10-027-632-224707 S	14.4 80.0 610 5 US-10-027-632-224706 S	14.4 80.0 610 5 US-10-02/-632-224/05 5	171 14.4 80.0 610 4 US-09-925-065A-462216 5	170 14.4 80.0 606 4 US-09-925-065A-451501 S	10 10 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
41 14 77.8 201 8 US-10-719-993-22356	14 77.8 201 7 US-10-741-601-7861	38 14 77.8 25 8 US-10-719-900-763654	37 14.4 80.0 3186778 6 US-10-027-632-174961	35 14.4 80.0 329019 8 US-10-388-838-48	234 14.4 80.0 169567 8 US-10-719-993-6774	232 14.4 80.0 142201 8 05-10-367-094-99 233 14.4 80.0 142976 7 US-10-367-094-99	231 14.4 80.0 127767 7 US-10-322-281-797	30 14.4 80.0 116704 5 US-10-087-192-1456	29 14.4 80.0:115218 8 US-10-278-698-769	228 14.4 80.0 115218 8 US-10-278-698-255	220 14.4 80.0 103747 7 US-10-450-826-46	225	24	223 14.4 80.0 8323 3 US-09-970-597-2	222 14.4 80.0 7036 5 US-10-178-782-3	221 14.4 80.0 3153 6 US-10-104-047-327	220 14.4 80.0 2754 3 US-09-938-842A-1786	219 14.4 80.0 2754 3 US-09-938-842A-1786	18 14.4 80.0 2621 10 US-11-097-143-36034 Sequen	15	215	216 177 80.0 1737 7 IS=00=525-66257 75133 8 Sequence	213 14.4 80.0 1262 4 US-09-26-06-07 273 Segments	212 14 4 80 0 1262 4 115-09-925-065A-727392 Sequence	212 14.4 80.0 1262 4 US-09-925-065A-727391 Sequence	11 14 4 80.0 939 8 US-10-926-684-403 Seguence	210 14 4 80 0 939 6 US-10-170-097-403 Sequence	200 14.4 80.0 878 6 US-10-027-632-136896 Sequence	207 14.4 80.0 878 K W2-10-027-632-136896 Segmenc	00.0 04.4 00.0 04.5 0 05-10-01-021-0213 3 360 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	05 14.4 80.0 845 6 HS-10-027-632-157312 Sequence	os 14 4 80 0 845 5 US-10-027-632-157313 Sequenc	03 14.4 00.0 013 7 00 03 323 0031 1220 Sequence	0. 0 0 0 12 4 170 000 000 000 000 000 000 000 000 000	01 14.4 00.0 013 4 00.00 001 -/	00 1 14 4 00 0 0 13 4 TEC-00-02E-05EA-7332A Semient	200 14 4 80 0 733 7 IV-10-424-599-136542 Sequence	199 14.4 RD 0 703 6 US-10-027-632-25471 Sequence	198 14.4 80.0 703 6 US-10-027-632-25470 Sequence	197 14.4 80.0 703 6 US-10-027-632-25469 Sequence	196 14.4 80.0 703 6 US-10-027-632-25468 Sequence	195 14.4 80.0 703 5 US-10-027-632-25471 Sequenc	194 14.4 80.0 703 5 US-10-027-632-25470 Sequence	193 14.4 80.0 703 5 US-10-027-632-25469 Sequence	192 14.4 80.0 703 5 US-10-027-632-25468 Sequence	14.4 80.0 663 6 US-10-027-632-227862 sequence	14.4 80.0 663 5 US-10-02/-632-22/862 Sequence	14.4 80.0 658 8 05-10-928-8028 303863 Seguente	14.4 80.0 658 6 US-10-17-10-2	14.4 80.0 636 8 05-10-33/-930-3040 Generalization	186 14.4 80.0 63.6 US+10-02/-632-2/3221 Sequence	: 185 14.4 80.0 632 6 US-10-02/-632-2/-5320 Sequence	184 14.4 80.0 632 5 US-10-02/-632-2/5521 Sequence	183 14.4 80.0 632 5 US-10-02/-632-2/-5320 Sequence	182 14.4 80.0 621 10 031 633 033 033 033 033	100 14:4 00:0 05:10:10:10:10:10:10:10:10:10:10:10:10:10:	101 14 4 00 0 616 6 116-027-633-307702 Segmenter	180 0 616 6 III-10-027-632-93222 Sequenc	: 179	: 178 14.4 80.0 616 5 US-10-027-632-93222 Sequenc	14.4 80.0 610 6 05-10-02/-02/-02/-03/-03/-03/-03/-03/-03/-03/-03/-03/-03	14 4 00 6 610 6 IIS-10-027-632-224707 Semienc	.76	75 14.4 80.0 610 6 US-10-027-632-224705 Sequence	74 14.4 80.0 610 5 US-10-027-632-224707 Sequence	14.4 80.0 610 5 US-10-027-632-224706 Sequence	14.4 80.0 610 5 US-10-02/-632-224/05 Sequence	171 14.4 80.0 610 4 US-09-925-065A-462218 Sequence	170 14.4 80.0 606 4 US-09-925-065A-451501 Seguenc	17. 00.0 606 4 170-026-06-4230 0 00 0 10 075
41 14 77.8 201 8 US-10-719-993-22356 Sequence 2235	14 77.8 201 7 US-10-741-601-7861 Sequence 7861	38	37 14.4 80.0 3186778 6 US-10-027-632-174961 Sequence 174	35	234 14.4 80.0 169567 8 US-10-719-993-6774 Sequence 67	233 14.4 80.0 142976 7 US-10-367-094-99 Sequence 99	231 14.4 80.0 127767 7 US-10-322-281-797 Sequence 79	30 14.4 80.0 116704 5 US-10-087-192-1456 Sequence 14	29 14.4 80.0 115218 8 US-10-278-698-769 Sequence 76	228 14.4 80.0 115218 8 US-10-278-698-255 Sequence 25	227 14 4 80.0 103747 7 US-10-450-826-46 Sequence 46.	225 14.4 80.0 71953 5 US-10-087-192-1057 Sequence 10	24	223 14.4 80.0 8323 3 US-09-970-597-2 Sequence 2,	222 14.4 80.0 7036 5 US-10-178-782-3 Sequence 3,	221 14.4 80.0 3153 6 US-10-104-047-327 Sequence 32	220 14.4 80.0 2754 3 US-09-938-842A-1786 Sequence 178	219 14.4 80.0 2754 3 US-09-938-842A-1786 Sequence 178	18 14 4 80.0 2621 10 US-11-097-143-36034 Sequence 30	15 14.4 80.0 2136 / US-10-424-599-136544 Sequence 130	215	214	213 14.4 80.0 1262 4 102-09-323-0634-727393 Sequence 72:	212 14 4 80 0 1262 4 IIS-09-925-065A-727392 Sequence 72:	212 14.4 80.0 1262 4 US-09-925-065A-727391 Sequence 72	11 14 4 80.0 939 8 US-10-926-684-403 Sequence 40	210 14 4 80 0 939 6 US-10-170-097-403 Sequence 400	200 14.4 80.0 878 6 US-10-027-632-136896 Sequence 130	20/ 14.4 80.0 843 5 TS-10-02/-532-136896 Sequence 130	07 14.4 00.0 043 0 00-10-10-1-157313 Sequence 15.	05 14.4 80.0 845 6 US-10-027-632-157312 Sequence 15.	os 14.4 80.0 845 5 US-10-027-632-157313 Sequence 15.	0.3 14.4 80.0 845 5 US-10-027-632-157312 Sequence 15.	02 14.4 00.0 013 4 05-09-941-0015-7414-7 00-01-01-01-07-7414-72-72-72-72-72-72-72-72-72-72-72-72-72-	OF 14.4 BULV BID 4 DECOMPANDE ACENTARY CONTROL 72	00	200 14.4 80.0 733 7 IIS-10-424-599-136542 Sequence 130	199 14 4 80.0 703 6 US-10-027-632-25471 Sequence 25	198 14.4 80.0 703 6 US-10-027-632-25470 Sequence 25	197 14.4 80.0 703 6 US-10-027-632-25469 Sequence 254	196 14.4 80.0 703 6 US-10-027-632-25468 Sequence 254	195 14.4 80.0 703 5 US-10-027-632-25471 Sequence 254	194 14.4 80.0 703 5 US-10-027-632-25470 Sequence 254	193 14.4 80.0 703 5 US-10-027-632-25469 Sequence 254	192 14.4 80.0 703 5 US-10-027-632-25468 Sequence 254	14.4 80.0 663 6 US-10-027-632-227862 sequence 22	14.4 80.0 663 5 US-10-02/-632-22/862 Sequence 22	10. 0 10. 10. 10. 10. 10. 10. 10. 10. 10	14.4 80.0 05.0 05.10-110-107-1404 05040-110-404	14.4 80.0 636 8 US-IU-39/-930-30463 CEMICENCE DO	186 14.4 80.0 632 6 US-10-02/-03/-1 Sequence C/-	185 14.4 80.0 632 6 US-10-02/-632-2/5520 Sequence 2/5	184 14.4 80.0 632 5 US-10-02/-632-2/5521 Sequence 2/3	183 14.4 80.0 632 5 US-10-02/-632-2/3520 Sequence 2/3	182 14.4 80.0 621 T 10.07 F143-30033 Commence of	10 10 11 14.4 00.0 010 10 110 110 12 12 12 12 12 12 12 12 12 12 12 12 12	160 14.4 00.0 016 6 Us-10-027-632-307702 Semience 307	180 14 4 80 0 616 6 US-10-027-632-93222 Sequence 932	: 179	: 178 14.4 80.0 616 5 US-10-027-632-93222 Sequence 93	14.4 00.0 otto 02110-021-032-02-03-03-03-03-03-03-03-03-03-03-03-03-03-	14 00 610 6 US-10-632-224707 Semience 224	.76	75 14.4 80.0 610 6 US-10-027-632-224705 Sequence 22	74 14.4 80.0 610 5 US-10-027-632-224707 Sequence 224	14.4 80.0 610 5 US-10-027-632-224706 sequence 22.	14.4 80.0 610 5 US-10-02/-632-224/05 Sequence 22:	171 14.4 80.0 610 4 US-09-925-065A-462218 Sequence 46	170 14.4 80.0 606 4 US-09-925-065A-451501 Sequence 450	17 and 17
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-925-065A-525294
-925-065A-727474
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Sequence 17868, A
Sequence 669, Appl
Sequence 10, Appl
Sequence 11, Appl
Sequence 5621, Ap
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Sequence 58, Appl
Sequence 58, Appl
Sequence 58, Appl
Sequence 58, Appl
Sequence 61, Appl
Sequence 615, Ap
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 INVENTION: KINASE SUPPRESSOR OF RAS INACTIVATION FOR THERAPY OF RAS MEDIAT
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RESULT 3
US-10-727-358-28/c
; Sequence 28, Application US/10727358
; Publication No. US20050037455A1
; GENERAL INFORMATION:
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APPLICANT: Kolesnick, Richard N.
APPLICANT: Xing, Hong-Mei R.
TITLE OF INVENTION: KINASE SUPPRESSOR OF RAS
TITLE OF INVENTION: TUMORICENESIS
FILE REFERENCE: 1216-1-006CIP
CURRENT APPLICATION NUMBER: US/10/727,358
CURRENT FILING DATE: 2003-12-03
PRIOR APPLICATION NUMBER: 60/384,228
PRIOR FILING DATE: 2002-05-30
PRIOR PPLICATION NUMBER: 60/460,023
PRIOR APPLICATION NUMBER: FOT/US03/16961
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR APPLICATION DATE: 2003-04-03
PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR FILING DATE: 2003-05-29
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; ORGANISM: Homo sapiens
US-10-727-358-5
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SOFTWARE: PatentIn version
SEQ ID NO 15
LENGTH: 18
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SEQ ID NO 5
LENGTH: 18
APPLICANT: Memorial Sloan-Kettering Cancer Center APPLICANT: Kolesnick, Richard N. APPLICANT: Xing, Hong-Mei R.
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NUMBER OF SEQ ID NOS: 38
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ORGANISM: Artificial Sequence
FEATURE:
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FILING DATE: 2003-12-03
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TITLE OF INVENTION:

1216-1-006CIP

TUMORIGENESIS

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Matches
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LENGTH: 25
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Publication No. US20050214823A1
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                                                                                                                                                                                        GENERAL INFORMATION:
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ORGANISM: Mus musculus
-11-036-317-881396
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             APPLICANT: Williams, Alan
APPLICANT: Blume, John
TITLE OF INVENTION: Method of Analysis of Alternative Splicing
FILE REFERENCE: 3854.1
CURRENT APPLICATION NUMBER: US/11/036,317
CURRENT FILING DATE: 2005-01-13
PRIOR APPLICATION NUMBER: US 60/536,639
PRIOR FILING DATE: 2004-01-13
                                                                                                                                                                                                                                                          11-036-317-904638
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NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
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CURRENT FILING DATE: 2003-12-03
CURRENT PELICATION NUMBER: 60/384,228
PRIOR APPLICATION NUMBER: 60/460,023
PRIOR APPLICATION NUMBER: 60/460,023
PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: PCT/US03/16961
                                                                                                                                                                                                            equence 904638, Application US/11036317 ublication No. US20050214823A1
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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PRIOR FILING DATE: 2004-01-13
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APPLICANT: Blume, Jo
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ITLE OF INVENTION: Method of Analysis of Alternative Splicing in
ILE REFERENCE: 3654.1
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ORGANISM: Artificial Sequence
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Local Similarity · 100.0%;
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SEQ ID NOS
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Pred. No. 27;
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US-11-036-317-904638
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US-11-036-317-910916
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APPLICANT: Williams,
APPLICANT: Blume, 30
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                                                                   SEQ ID NO 1
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Best Local
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CURRENT APPLICATION NUMBER: US/11/036,317
CURRENT FILING DATE: 2005-01-13
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                                                                                                                   PRIOR APPLICATION NUMBER: 60/384,228
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: 60/460,023
PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR FILING DATE: 2003-05-29
                                                                                                                                                                                                                                                                               APPLICANT: Kolesnick, Richard N.
APPLICANT: Xing, Hong-Mei R.
TITLE OF INVENTION: KINASE SUPPRESSOR OF RAS INACTIVATION FOR THERAPY OF RAS MEDIATE
TITLE OF INVENTION: TUMORIGENESIS
FILE REFERENCE: 1216-1-006CIP
                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/727,358
CURRENT FILING DATE: 2003-12-03
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TYPE: DNA ORGANISM: Homo sapiens
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ITLE OF INVENTION: Method of Analysis of Alternative Splicing in Mouse
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Pred. No.
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; ORGANISM: Homo sapiens
US-10-727-358-24
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US-10-727-358-25
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                                                                               SOFTWARE: Pa
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LENGTH: 121
TYPE: DNA
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Publication No. US20050037455A1
GENERAL INFORMATION:
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                                                                                                                                             PRIOR APPLICATION NUMBER: 60/384,228
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: 60/460,023
PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR FILING DATE: 2003-05-29
                                                                                                                                                                                                                                                               APPLICANT: Xing, Hong-Mei R.

TITLE OF INVENTION: KINASE SUPPRESSOR OF RAS INACTIVATION FOR THERAPY OF RAS MEDIATED TITLE OF INVENTION: TUMORIGENESIS

FILE REFERENCE: 1216-1-006CIP

CURRENT APPLICATION NUMBER: US/10/727,358

CURRENT FILLING DATE: 2003-12-03
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PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: 60/460,023
PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR FILING DATE: 2003-05-29
NUMBER OF SEQ ID NOS: 38
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TITLE OF INVENTION: KINASE SUPPRESSOR OF RAS INACTIVATION FOR THERAPY OF RAS MEDIATER
TITLE OF INVENTION: TUMORIGENESIS
FILE REFERENCE: 1216-1-006CIP
CURRENT APPLICATION NUMBER: US/10/727,358
CURRENT FILING DATE: 2003-12-03
                                                                                                                                  NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Memorial Sloan-Kettering Cancer Center APPLICANT: Kolesnick, Richard N. APPLICANT: Xing, Hong-Mei R.
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APPLICANT: Xing, Hong-Mei R.
                                              TYPE: DNA
                                                               ENGTH:
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Local Similarity 100.0%; (
nes 18; Conservative 0;
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Pred. No.
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Pred. No. 22;
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Query Match

100.0%; Score 18;

DB 8;

Length 2601;

RESULT 12 US-10-727-358-11

Sequence 11, Application US/10727358 Publication No. US20050037455A1

GENERAL INFORMATION:

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                                                                                                                              ; TYPE: DNA; ORGANISM: Mus musculus US-10-085-117-32
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US-10-085-117-32
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US-10-085-117-33
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                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 32
                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 32, Applica Publication No. US20 GENERAL INFORMATION:
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                                                                                Query Match
Best Local Similarity
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Matches 18; Conser
                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 09/798,586
PRIOR FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 361
                                                                                                                                                                                                                                                                                                                      APPLICANT: Engelhard, Bric K.
TITLE OF INVENTION: NOVEL COMPOSITIONS AND
FILE REFERENCE: 529452000121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: MOTTIS, David W.
APPLICANT: Engelhard, Eric K.
APPLICANT: Engelhard, Eric K.
TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER
FILE REFERENCE: 529452000121
CURRENT APPLICATION NUMBER: US/10/085,117
CURRENT FILING DATE: 2002-02-27
PRIOR APPLICATION NUMBER: US 09/798,586
PRIOR FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 361
                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/085,117
CURRENT FILING DATE: 2002-02-27
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Morris
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                                                                                                                                                                                  LENGTH: 4094
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296 CGGACCCTAGAGGCAAAG 313
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                              CGGACCCTAGAGGCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                Application US/10085117
No. US20030232334A1
                                                         100.0%; Score 18; DE nilarity 100.0%; Pred. No. 15; Conservative 0; Mismatches
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                                                                                                Length 4094;
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                                                                  Indels
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RESULT 14
US-11-036-317-904637
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US-11-036-317-881395
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PRIOR APPLICATION NUMBER: US 60/536,639

PRIOR FILING DATE: 2004-01-13

NUMBER OF SEQ ID NOS: 991174

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 881395
                                                                                                                                    Sequence 904637, Application US/11036317
Publication No. US20050214823A1
GENERAL INFORMATION:
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Best Local Similarity
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SEQ ID NO 11
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TYPE: DNA
ORGANISM: Mus musculus
-10-727-358-11
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PRIOR TILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: PCT/US03/16961
PRIOR FILING DATE: 2003-05-29
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CURRENT APPLICATION NUMBER: US/10/727,358
CURRENT FILING DATE: 2003-12-03
PRIOR APPLICATION NUMBER: 60/384,228
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-05-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 25
TYPE: DNA
                                                                           APPLICANT: Williams, Alan
APPLICANT: Blume, John
TITLE OF INVENTION: Method of Analysis of Alternative Splicing in Mouse
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                    CURRENT APPLICATION NUMBER: US/11/036,317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Blume, John
IITLE OF INVENTION: Method of Analysis of Alternative Splicing in Mouse
FILE REFERENCE: 3654.1
                                                          ILE REFERENCE: 3654.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PPLICANT: Xing, Hong-Mei R.
ITLE OF INVENTION: KINASE SUPPRESSOR OF RAS INACTIVATION FOR THERAPY OF RAS MEDIATEDITED OF INVENTION: TUMORIGENESIS
                                                                                                                                                                                                                                                                                                                                                              Local Similarity
mes 17; Conservat
APPLICATION NUMBER:
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94.4%;
US 60/536,639
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Pred. No. 1.
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Pred. No.
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US-11-036-317-904637
                                                                       CURRENT APPLICATION NUMBER: US/10/727,358
CURRENT FILING DATE: 2003-12-03
PRIOR APPLICATION NUMBER: 60/384,228
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: 60/460,023
PRIOR APPLICATION NUMBER: 60/460,023
PRIOR FILING DATE: 2003-04-03
PRIOR FILING DATE: 2003-05-29
PRIOR FILING DATE: 2003-05-29
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Sequence 910915, Application US/11036317

Publication No. US20050214823A1

GENERAL INFORMATION:

APPLICANT: Williams, Alan

APPLICANT: Blume, John

APPLICANT: Blume, John

APPLICANT: Muchams, Method of Analysis of Alternative Splicing in Mouse
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                     SEQ ID NO 8
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NUMBER OF SEQ ID NOS: 991174
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 3654.1
CURRENT APPLICATION NUMBER: US/11/036,317
CURRENT FILING DATE: 2005-01-13
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                     APPLICANT: Memorial Sloan-Kettering Cancer Center
APPLICANT: Kolesnick, Richard N.
APPLICANT: Xing, Hong-Mei R.
TITLE OF INVENTION: KINASE SUPPRESSOR OF RAS INACTIVATION FOR THERAPY OF RAS MEDIATE
TITLE OF INVENTION: TUMORIGENESIS
FILE REFERENCE: 1216-1-006CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/536,639 PRIOR FILING DATE: 2004-01-13
                                     SOFTWARE: PatentIn version 3.1
                                                        NUMBER OF SEQ ID NOS:
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17; Conserv
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94.48;
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Pred. No. 1
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Pred. No. 1
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ORGANISM: Artificial Sequence

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US-11-036-317-928166
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                                                                                                                SEQ ID NO 12378
LENGTH: 233
TYPE: DNA
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                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Compositions and Methods Relating to Osteoarthritis
FILE REFERENCE: 4231/2005
CURRENT APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR TILING DATE: 2001-03-12
                                                                                                                                                                                NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn ve
                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 991174
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/11/036,317
CURRENT FILING DATE: 2005-01-13
PRIOR APPLICATION NUMBER: US 60/536,639
PRIOR FILING DATE: 2004-01-13
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APPLICANT: Blume, John
TITLE OF INVENTION: Method of Analysis of Alternative Splicing in Mouse
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APPLICANT: Liew, C.C.
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                         ORGANISM: Human
FEATURE:
NAME/KEY: misc_feature
LOCATION: (13)...(13)
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OTHER INFORMATION: antisense oligonucleotide
    LOCATION: (13)..(13)
OTHER INFORMATION: n is a,
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                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 60/271,955
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; OTHER INFORMATION: n is a, US-10-085-783A-12378
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; LOCATION: (66)...(66)
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US-10-242-535A-12378
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 12378
LENGTH: 233
Type: No.
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Query Match
Best Local Similarity
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Compositions and Methods Relatiing to Osteoarthritis FILE REFERENCE: 4231/2002 CURRENT APPLICATION NUMBER: US/10/085,783A CURRENT FILING DATE: 2002-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR FILING DATE: 2001-02-28
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LOCATION: (66)..(66)
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OTHER INFORMATION: n is
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LOCATION: (39)..(39)
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OTHER INFORMATION: n is
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LOCATION: (39)..(39)
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LOCATION: (28)..(28)
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SEQ ID NO 522
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Best Local
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Patent No. US20020102531A1
SENERAL INFORMATION:
APPLICANT: HOrrigan, Stephen
TITLE OF INVENTION: Cancer Gene Determination
TITLE OF INVENTION: Sets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Homo sapiens
-09-964-824A-522
                                                                                                                                 LENGTH: 376
TYPE: DNA
ORGANISM: Homo sapiens
-09-969-708-67
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: US/60/237,608
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: US/60/237,425
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CURRENT FILING DATE: 2001-09-27
CURRENT FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 658
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MITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
MITLE OF INVENTION: Sets
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                                                                  Local Similarity
es 16; Conserv
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                                    GGACCCTAGAGGCAAAG 18
   GGACCCTAGAGGCAGAG 91
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94.1%;
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94.1%;
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Pred. No. 4.4e.
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Pred. No. 4.4e+02;
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                                                                                              CURRENT APPLICATION NUMBER: US/10/843,641A
CURRENT FILING DATE: 2004-05-12
PRIOR APPLICATION NUMBER: US/09/873,367
PRIOR FILING DATE: 2001-66-05
PRIOR APPLICATION NUMBER: US/09/954,531
PRIOR PILING DATE: 2001-09-18
PRIOR PILING DATE: 2001-09-18
PRIOR PILING DATE: 2001-09-25
PRIOR FILING DATE: 2001-09-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US/09/873,367 PRIOR FILING DATE: 2001-06-05
                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
TITLE OF INVENTION: Signature Gene Sets
FILE REPERENCE: 689290-189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                   APPLICANT: Avalon Pharmaceuticals, Inc.
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IITLE OF INVENTION: Signature Gene Sets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/954,456 FILING DATE: 2001-09-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/954,531 FILING DATE: 2001-09-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/962,832
                                                        APPLICATION NUMBER: US/09/962,436 FILING DATE: 2001-09-25
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16; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/969,708
                                     APPLICATION NUMBER: US/09/962,832
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94.1%;
US/09/964,824
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Pred. No. 4.4e
0; Mismatches
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RESULT 25
US-10-027-632-141547
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Sequence 141547, Application US/10027632

Publication No. US20020198371A1

GENERAL INFORMATION:
APPLICANT: Wang, David G.

APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129

CURRENT APPLICATION NUMBER: US/10/027,632

CURRENT FILING DATE: 2002-04-30

PRIOR APPLICATION NUMBER: US 60/218,006

PRIOR APPLICATION NUMBER: US 60/198,676
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Best Local S
Matches 16
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SEQ ID NO 110504
LENGTH: 383
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants FILE REFERENCE: 38-21(5322) B CURRENT APPLICATION NUMBER: US/10/425,115 CURRENT FILING DATE: 2003-04-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/09/967,768
PRIOR FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: US/09/968,007
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NUMBER OF SEQ ID NOS: 8447
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Clone ID: MRT4577_32272C.1
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APPLICATION NUMBER: US/09/969,347
FILING DATE: 2001-10-02
APPLICATION NUMBER: US/09/969,708
FILING DATE: 2001-10-03
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Similarity 94.1%;
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Cao, Yongwei
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94.1%;
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Pred. No. 4.4e+02;
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US-10-027-632-141547
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PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
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Best Local (
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SEQ ID NO 141547
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NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FRENSEQ for Windows Version 4.0
SEQ ID NO 141547
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CORMERCE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-04-30
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
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                                                                                                                                                                           TYPE: DNA ORGANISM: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/198,676
                                                                                                                                                                                                                 LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/167,363
FILING DATE: 1999-11-23
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255 GGÁCCCCÁGAGGCÁAÁG
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                                     2 GGACCCTAGAGGCAAAG 18
                                                                                                Similarity
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                                                                             Conservative
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                                                                                              Score 15.4; DB 6;
Pred. No. 4.2e+02;
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Pred. No. 4.2e+02;
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RESULT 27 US-09-764-872-162/c

Sequence 162, Application US/09764872

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RESULT 28
US-10-027-632-202268
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                                                                         RESULT 29
US-10-027-632-202268
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; ORGANISM: Human
US-10-027-632-202268
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; ORGANISM: Homo sapiens
US-09-764-872-162
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FASTSEQ FOr Windows Version 4.0
SEQ ID NO 202268
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CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper
NUMBER OF SEQ ID NOS: 957
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 162
LENGTH: 514
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              Sequence 202268, Application US/10027632 Publication No. US20030204075A9 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Publication No. US20
GENERAL INFORMATION:
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
FILE REFERENCE: 10827.129
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TITLE OF INVENTION: Nucleic Acids, Proteins, and
FILE REFERENCE: PA125
APPLICANT: Wang, David G.
                                                                                                                                                                                                                                                                                                                                                                      ENGTH: 546
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                                                                                                                                                                                                                                                      Local
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/18,676
PRIOR PILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-03-29
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
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APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
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CURRENT FILING DATE: 2003-04-28
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PRIOR FILING DATE: 2000-02-24
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ACCESSION VERSION KEYWORDS

BB629501

SOURCE

Mus musculus (house mouse) BB629501.1 GI:15399640 LOCUS DEFINITION RESULT 1 BB629501

BB629501
BB629501 RIKEN full-length enriched, adult cDNA clone 9830134F20 5', mRNA sequence.

male

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REFERENCE
AUTHORS
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Contact: Yoshihide Hayashizaki
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Arakawa, T., Carninci, P.,
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The Institute of Physical and Chemical Research (RIKEN)
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/note="Site_1: Sall; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Rike
                                                                                                                                                                                                                                                                                                                 Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken
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Sakazume, N., Sato, K., Shiraki, T., Waki, K., Kawai, J., Aizawa, K.,

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Sandorn, R., Lander, E.S.,

Analysis, of the mouse transcriptome based on functional annotation
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Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs
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                                                                                                                                                                                                                                                                                                                                                                                  Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Jan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nature 420, 563-573
12466851
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
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EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Yoshihide Hayashizaki
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Watahiki, A., Muramatsu, M. and Hayashizaki, Y.
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                                                                                                                                                                                                                                                                                                                                                                                      Yokohama, Kanagawa 230-0045, Japan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
1 (bases 1 to 688)
                                                                                                                                                                                           BB613725 RIKEN full-length enriched, 0 day neonate musculus cDNA clone 4831413M17 5', mRNA sequence. BB613725
                                                                                                                                                                                                                                                                                                                                                                                                                           1 CGGACCCTAGAGGCAAAG 18
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Please visit our web site (http://genome.gsc.riken.go.jp) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequencing pipeline with 384 multicapillary sequencer. Genome 10 (11), 1757-1771 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genes.
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                                                                                                    Mus musculus
                                                                                                                                                                     BB613725.1 GI:15395652
                                                                                                                        Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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RIKEN integrated sequence analysis (RISA) system--384-format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Site 1: Sall; Site 2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was primed with a primer [5' GAGARAGARA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /dev_stage="6 days neonate'
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lib="RIKEN full-length enriched, 6 days neonate
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ches 0;
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                                                                                 Euteleostomi;
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Carninci, P., Fukuda, S.,

Furuno, M., Hanagaki, T.,

COMMENT TITLE JOURNAL

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source
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The Institute of Physical and Chemical Research (RIKEN)
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                                     prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken
                                                                                                                                                                                                               cissues
Vector: a modified pBluescript KS(+) after bulk excision
from Lambda FT.C T "
                                                                                                                                                                                                                                                                                                                                                                                                      /note="Site_1: Sall; Site_2: BamHI; cDNA library was
                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone_lib="RIKEN full-length enriched, 0 day neonate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'dev_stage="0 day neonate"
'lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sex="mixed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        db_xref="taxon:10090"
clone="4831413M17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Mus musculus"
mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tissue_type="head"
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FEATURES

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                                          Adachi J., Aizawa, K., Akimura, T., Arakawa, T., Bono, H., Carninci, F. Fukuda, S., Furuno, M., Hanagaki, T., Hara, A., Hashizune, W., Hayashida, K., Hayatsu, N., Hiramoto, K., Hiraoka, T., Hirozane, T., Hori, F., Imotani, K., Ishii, Y., Itoh, M., Kagawa, I., Kasukawa, T., Koya, S., Kurihara, C., Matsuyama, T., Kondo, S., Konno, H., Kouda, M., Koya, S., Kurihara, C., Matsuyama, T., Myazaki, A., Murata, M., Nakamura, M., Nishi, K., Nomura, K., Numazaki, R., Ohno, M., Ohsato, N., Okazaki, Y., Saito, R., Saitoh, H., Sakai, C., Sakai, K., Sakazume, N., Sano, H., Sasaki, D., Shibata, K., Shinagawa, A., Shiraki, T., Sogabe, Y., Tagami, M., Tagawa, A., Shiraki, T., Takaku-Akahira, S., Takeda, Y., Tanaka, T., Tomaru, A., Toya, T., Yasunishi, A., Muramatsu, M., and Hayashizaki, Y.
                                                                                                                                                                                                                                                                                                                                                                 Group Phase I & II Team.
Analysis of the mouse transcriptome based on of 60,770 full-length cDNAs
Nature 42b, 563-573 (2002)
6 (bases 1 to 2751)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shibata, K., Itoh, M., Aizawa, K., Nagaoka, S., Sasaki, N., Carninci, P., Konno, H., Akiyama, J., Nishi, K., Kitsunai, T., Tashiro, H., Itoh, M., Sumi, N., Ishii, Y., Nakamura, S., Hazama, M., Nishine, T., Harada, A., Yamamoto, R., Matsumoto, H., Sakaguchi, S., Ikegami, T., Kashiwagi, K., Fujiwake, S., Inoue, K., Togawa, Y., Izawa, M., Ohara, B., Watahiki, M., Yoneda, Y., Ishikawa, T., Ozawa, K., Tanaka, T., Matsuura, S., Kawai, J., Okazaki, Y., Muramatsu, M., Inoue, Y., Kira, A. and Hayashizaki, Y. RIKEN integrated sequence analysis (RISA) system-384-format sequencing pipeline with 384 multicapillary sequencer Genome Res. 10 (11), 1757-1771 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y. Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genome Res. 10 (10), 1617-1630 (2000)
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Submitted (16-JUL-2001) Yoshihide Hayashizaki, The Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning
Meth. Enzymol. 303, 19-44 (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The FANTOM Consortium and the RIKEN Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The RIKEN Genome Exploration Research Group Phase II Team and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AK036566.1 GI:26085304
HTC; CAP trapper.
Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AK036566 2751 bp mRNA linear HTC 03-APR-2004 Mus musculus adult male bone cDNA, RIKEN full-length enriched library, clone:9830134F20 product:kinase suppressor of ras, full
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nature 409, 685-690 (2001)
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Mammalia, Eutheria, Euarchontoglires, Glires, Rodentia,
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Pred. No.
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Physical and Chemical Research (RIKEN), Laboratory for Genome

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zmrww005_0A10-002-b11.80 zmrww005 Zea mays cDNA 5',
CK370327-
CK370327.1 GI:40336257
                                                                                                                                                                                                      Contact: Hans Bohnert
University of Illinois, Urbana-Champaign
1201 West Gregory Drive, Urbana, IL 61801
Tel: 217-265-5475
Fax: 217-333-5574
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zea mays
Zea mays
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                                                                                                                                                                                                                                                                                                                            Bohnert, H., Sharp, R.E., Springer, G.K., Poroyko, V., Fredricksen, M., Sharp, L.G., Spollen, W.G., Ries, J., Guillen, A., Khambati, A., Topinka, C., Davis, G.E., Schachtman, D., Wu, Y. and Nguyen, H.T. NSF Grant DBI-0211842: Functional Genomics of Root Growth and Root Signaling Under Drought Unpublished (2003)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     URL:http://genome.gsc.riken.jp/
URL:http://fantom.gsc.riken.jp/.
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Please visit our web site for further details.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clade; Panicoideae; Andropogoneae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
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                                                                                                                                                                                          bohnerth@life.uiuc.edu
/organism="Zea mays"
/mol_type="mRNA"
/db_xref="texon:4577"
/clone_lib="zmrww005"
/note="Library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="kinase suppressor of ras (MGD|MGI:105051, GB|NM_013571, evidence: BLASTN, 100%, match=1985)"
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/clone_lib="RIKEN full-length enriched mouse cDNA library"
/dev_stage="adult"
                                                                                                                     Location/Qualifiers
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/db_xref="taxon:10090"
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/strain="C57BL/6J"
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    Library zmrww005 consists of the same cDNA
                                                                                                                                                                                                                                                               Urbana, IL 61801, USA
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sequenced from the S prime end. The sequence identifier uses the .e0' suffix because the library tag was at the 3 prime end and thus not identified. Samples were collected in Robert B. Shary's lab (University C.) Missouri-Columbial to construct three normalized CNNA libraries. Dark'sgrown maize seedlings with primary roos for the sequence identified. Samples were collected in Robert B. Shary's lab (University C.) Missouri-Columbial to construct three normalized CNNA libraries. Dark'sgrown maize seedlings with primary roos for the root and the root appears and root water poemical (1-1.6 MHz) very pential libraries are from the pinetro of the root apex and root a root seed for a certain this 500 vacts were constructed in the pinetro of the root apex and root a root method (2). The three normalized CNNA libraries were constructed in the lab of Hans Bohnert (1) and ed. 1997). This method worked in the segment (2). The three normalized CNNA libraries were constructed in the lab of Hans Bohnert (1) and ed. 1997). This method worked in the root pex and root a principle of the root was added to the order than the root pex and root and the root in the lab of Hans Bohnert (1) and ed. 1997). This method worked in the root pex and root primers were constructed in the lab of Hans Bohnert (1) and ed. 1997). This method worked in the root appears the root and the root and the root was added to the root was added to the state of the modified Oligo(dT) primers. The three normalized CNNA pagnets of the root was added to the standard to double-stranded CNNA and tagged by using modified Oligo(dT) primers. One of 4 sequence tags added to each sequence tags and the root and root and the root of the root was added to the stranded clonas were adapticed (1400 bp). Site of the root was added to the stranded clonas were adapticed (1400 bp) in the primary libraries before amplification the root was a root of the modified Oligo(dT) primers. The cDNA was elected to double-strandard into bearth the sequence tag and the formal root of the roo

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REFERENCE
AUTHORS
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Unpublished (2003)
Contact: Hans Bohnert
University of Illinois, Urbana-Champaign
1201 West Gregory Drive, Urbana, IL 61801, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     zmrww00_0A10-002-b11.s0
CF633642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CF633642.1 GI:37392777
                                                                                                         1,000 roots were used for each of the low water potential libraries (zmrws05 and zmrws48) while 500 roots were combined from each of the two time points at high water potential (zmrww00). Each root was divided into 4 segments (distances are from the junction of the root apex and root cap): segment 1, 0-3 mm plus the root cap; segment 2, 3-7 mm; segment 3, 7-12 mm; segment 4, 12-20 mm. (For detatils of conditions see (1) with nutrient modifications as in (2)). The three normalized cDNA libraries were constructed in the lab of Hans Bohnert (University of Illinois-UC). Total RNA was extracted by the 'hot Phenol' method (Plant Molecular Biology manual. D5: 1-13, 2nd ed., 1997). This method worked in eliminating carbohydrate material present in the root tips. The integrity of the RNA was verified by denaturing agarose gels and spectrophotometry (ratio A260/280). Poly(A)+mRNA was isolated twice from total RNA was converted to double tranship and tractions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bohnerth@life.uiuc.edu
was converted to double-stranded cDNA and tagged by using modified Oligo(dT) primers. One of 4 sequence tags corresponding to a different segment of the root was added to the 3'-end of the modified Oligo(dT) primers, including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         normalized cDNA libraries. Dark-grown maize seedlings with primary roots 12-20 mm in length were transplanted to high (-0.03 MPa) or low water potential (-1.6 MPa) vermiculite, and harvested at 5 h and 48 h after transplanting. About
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="zmrww00"
/note="Samples were collected in Robert E. Sharp's lab
/note=sity of Missouri-Columbia) to construct three
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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TAG_TISSUE=Not found
TAG_SEQ=Not found"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                _type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91.1%;
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Pred. No. 1.2e+03;
0; Mismatches 1;
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Zea mays cDNA 3', mRNA sequence.
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the root was added primers, including
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1 CGGACCCTAGAGGCAAAG 18
                                                                                                                                                                                                                      n 91.1%;
Similarity 94.4%;
                                                                                                                                                                                 Conservative
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Non-bybridized single-stranded DNA circles were separated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Non-hybridized single-stranded DNA circles were separated from hybridized DNA rendezed partially double-stranded and electroporated into DH10B. The total number of clones with insert was: zmrws05: 2.0x107; zmrws48: 4.2x107; zmrws00: 1.1x107. The background of empty clones was less than 2%. Insert size, determined by PCR of the entire library, ranged from 0.5kb to 2.5kb. (1) Sharp R E; Silk W K; Hsiao T C. Growth of the Maize Primary Root at Low Water Potentials I. Spatial Distribution of Expansive Growth. Plant Physiology (Rockville). 87(1). 1988. 50-57. (2) Spollen W G; LeNoble M E; Samuels T D; Bernstein N; Sharp R E. Abscisic acid accumulation maintains maize primary root elongation at low water potentials by restricting enotype the production. Plant Physiology (Rockville). 122(3).
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TAG_SEQ=Not found"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000. 967-976
272
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                                                                                                                                                                                                                 Score 16.4; DB 6;
Pred. No. 1.2e+03;
                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                      Length 370;
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Query Match Best Local S Matches 17

RESULT 7

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DEFINITION ACCESSION BE717998/c LOCUS **KEYWORDS** VERSION 910 bp mRNA linear BE717998 BE717998 GI:10106263

mRNA sequence.

ORGANISM Homo sapiens (human) Eukaryota; EST

1 (bases 1 to 510)
Dias Neto, B., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deoliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and Hominidae; Mammalia; ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; | Butheria; Buarchontoglires; Primates; Catarrhini; Homo.

REFERENCE

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REFERENCE
AUTHORS
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AG354283
                                                                                                                               REFERENCE
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KEYWORDS
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PUBMED
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17;
                                                                                                                                                                    Contribution of Asian mouse subspecies Mus musculus molossinus genomic constitution of strain C57BL/6J, as defined by BAC-end sequence-SNP analysis
Genome Res. 14 (12), 2439-2447 (2004)
                                                                                                                                                                                                                                                                                                                                                             Mus musculus molossinus (Japanese wild mouse)
Mus musculus molossinus
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Eut
Mammalia; Butheria; Euarchontoglires; Glires; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                              Abe, K., Noguchi, H., Tagawa, K., Yuzuriha, M., Ezawa, K., Saitou, N., Hattori, M., Sakaki, Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence start: 20 High quality sequence stop: 429.
and Chemical Research (RIKEN), Genomic Sciences Center (GSC);
1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa, 230-0045, Japan
(E-mail:hattori@gsc.riken.jp, URL:http://hgp.gsc.riken.go.jp/,
                                                                                                                                                                                                                                                            Shiroishi, T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AG354283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus molossinus DNA,
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600-001-a09&t3=2000-06-25&t4=1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: +55-11-2704922
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                                                                 Submitted (17-NOV-2003) Masahira Hattori, The Institute of Physical
                                                                                       Direct Submission
                                                                                                       2 (bases 1 to 704)
Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y.
                                                                                                                                                                                                                                                                                                                                           Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AG354283.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CGGACCCTAGAGGCAAAG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=Torgan: head neck; Vector: puc18; Site 1: SmaI; Site 2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cissue mRNA and cDNA amplification were performed under
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone_lib="HT0790"
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Pred. No. 1.2e+03;
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clone:MSMg01-152G07.T7,
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                                                                                                                                                                                                                                                                                                                                                                                       Euteleostomi;
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R.Site 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       744 bp mRNA linear EST 11 AGENCOURT 55789292 NICHD_XGC_FAB Xenopus laevis cDNA clone IMAGE:8069448 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               library availability, please contact Kuniya Abe (abe@rtc.rike Tsukuba Institude, Bio Resource Center, The Institute of Physical and Chemical Research (RIKEN) 3-1-1 Koyadai, Tsukuba, 305-0074 Japan phone, 81-298-36-9189, fax: 81-298-36-9199
                                                                                                                                        CDNA Library Preparation: Express Genomics CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Agencourt Bioscience Corporation Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM17342 row: d column: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xenopus laevis (African clawed frog)
Xenopus laevis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Clones are derived from the mouse BAC library MSMg01. For BAC library availability, please contact Kuniya Abe (abe@rtc.riken.jp).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel:81-45-503-9111,
                                                                                                   High quality sequence stop: 623
Location/Qualifiers
                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Tom Sargent
                                                                                                                                                                                                                                                                                                                                    National Cancer Institute / NIH
Bldg. 31 Rm10A07 Bethesda, MD 20892
                                                                                                                                                                                                                                                                                                                                                                        Contact: Daniela S. Gerhard, Ph.D. Office of Cancer Genomics
                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kenopodinae; Kenopus; Kenopus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CGGAACCTAGAGGCAAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 744)
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/mol_type="genomic DNA"
/sub species="molossinus"
/db_xref="taxon:57486"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="mixture of
/clone_lib="MSMg01 Mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="MSMg01-152G07.T7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                     organism="Xenopus laevis"
/mol_type="mRNA"
                      db xref="taxon:8355"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sex="male
clone="IMAGE:8069448"
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: EcoRI
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94.4%;
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Pred. No. 1.3e+03;
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Male BAC Library"
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CX910994
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2004)
Other ESTB: JGI CAAN2676.rev
Contact: Lindquist,E.A., Richardson,P.
DOE Joint Genome Institute
2800 Mitchell Drive, Walnut Creek, CA 94598, USA
                                                                                                                                                                                                                                                                                                                                                                                          Naming Conventions: EST name is generated by the concatenation of the JGI Clone Id and the direction of sequencing. The suffix '.fwd' indicates a forward sequencing read of the insert. It does not necessarily reflect the orientation of the insert.

Plate: CAAN 0025 row: h column: 22
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Xenopus tropicalis
Xenopus tropicalis
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopudinae; Xenopus; Silurana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CX910994

JGI_CAAN2676.fwd NIH_XGC_tropTe4 Xenopus tropicalis cDNA clone
IMAGE:7688136 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA Sequencing: DOE Joint Genome Institute: http://www.jgi.doe.gov
Clone Distribution: I.M.A.G.E. Consortium/LLNL:
http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA Library Preparation: DOE Joint Genome Institute: http://www.jgi.doe.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cdna@jgi-psf.org
Tissue Procurement: Timothy Grammer (Richard M. Harland Laboratory,
University of California, Berkeley:
http://tropicalis.berkeley.edu/home)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Richardson, P., Lucas, S., Rokhsar, D., Detter, J.C., Ng, D.C., Brokstein, P. and Lindquist, E.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CGGACCCTAGAGGCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 925 296 5600
Fax: 925 296 5710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOE Joint Genome
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                                                                                                                                                                                                                                                                                                                                               quality sequence stop:
Location/Qualifiers
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/clone_Torgan: fat body; Vector: pExpress-1; Site_1: EcoRV;
Site_2: NotI; cDNA was primed using oligo-dT primer:
5'-pGACTAGTTCTAGATCGCGAGCGGCCGCCC(T)25-3' and cloned into
the EcoRV/NotI sites of pExpress-1. Size-selection 1.2kb
resulted in an average insert size of 1.8kb. This is a
primary library (normalized library is NICHD_XGC_FabN) and
was constructed by Express Genomics (Frederick, MD). Note:
this is a (http://xgc.nci.nih.gov/) Xenopus Gene
Collection library."
/clone lib="NIH XGC tropTe4"
/clone lib="NIH XGC tropTe4"
/note="Vector: pCMV5port6; Site_1: SalI; Site_2: NotI;
/note="Vector: pCMV5port6; Site_1: SalI; Site_2: NotI;
/note="Vector: pCMV5port6; Site_1: SalI; Site_2: NotI;
                                                                                                /dev_stage="Adult"
/lab_host="ElectroMAX DH10B"
                                                                                                                                                                                                                                                                                         organism="Xenopus tropicalis"
                                                                                                                                                                                                                                                                                                                           L. .817
                                                                                                                                                      tissue_type="Testes"
                                                                                                                                                                                        sex="male"
                                                                                                                                                                                                               clone="IMAGE: 7688136"
                                                                                                                                                                                                                                   xref="taxon:8364"
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94.4%;
                                                                                                                                                                                                                                                                     type="mRNA
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Pred. No. 1.3e+03;
0; Mismatches 1
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ORIGIN

FEATURES

Query Match Best Local Matches

1 Similarity 17; Conser

Conservative

91.1%;

Score 16.4; DB 8; Pred. No. 1.3e+03; 0; Mismatches 1;

Length 875;

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Gaps

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ORGANISM
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VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: Capabbe-remail.nih.gov
Tissue Procurement: Tom Sargent
cDNA Library Preparation: Express Genomics
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAMI7342 row: d column: 22
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1 (Dases 1 to 875)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                           High quality sequence start: 47: High quality sequence stop: 795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              National Cancer Institute / NIH
Bldg. 31 Rm10A07 Bethesda, MD 20892
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Daniela S. Gerhard, Ph.D. Office of Cancer Genomics
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Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xenopus laevis
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               /clone_lib="NICHD_XGC_FaB"
/rotes="Organ: fat body; Vector: pExpress-1; Site_1: EcoRV;
Site_2: Not1; cDNA was primed using oligo-dT primer:
5'-pGACTAGTTCTAGATCGCGAGCGGCCCCC(T)25-3' and cloned into
the EcoRV/Not1 sites of pExpress-1. Size-selection 1.2kb
resulted in an average insert size of 1.8kb. This is a
primary library (normalized library is NICHD_XGC_FabN) and
was constructed by Express Genomics (Frederick, MD). Note:
this is a (http://xgc.nci.nih.gov/) Xenopus Gene
Collection library."
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                                                                                                                                                                                                                                                           'clone="IMAGE:8069448"
'lab_host="DH10B TonA"
                                                                                                                                                                                                                                                                                                        organism="Xenopus laevis"
/mol_type="mRNA"
/db_xref="taxon:8355"
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Pred. No. 1.3e+03;
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RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Okazaki,Y. and Hayashizaki,Y.
Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)
Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning. Methods Enzymol. 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laboratory for Genome Exploration Research Group, RIKEN Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-
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RIKEN Mouse ESTs (Konno,H., et al.)
Unpublished (2000)
Contact: Yoshihide Hayashizaki
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         visit our web site (http://genome.rtc.riken.go.jp) for
prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken
                                                                                                                                                                                                                                                /tissue_type="cerebellum"
/dev_stage="7 days neonate"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Mus musculus"
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                                                                                                                                                                                                               clone_lib="RIKEN full-length enriched,
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                                                                                                                                        1: SalI; Site_2: BamHI; cDNA library was
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Dias Neto, B., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G. H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., Brunstein, A., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
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BE768604
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Project. This entry can be seen
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Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Simpson A.J.G
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/note="Organ: prostate tumor; Vector: pucl8; Site_1: SmaI; Site_2: SmaI; A mini-lībrary was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under
                                                                                                                                                                        /db_xref="taxon:9606"
/dev_stage="Adult"
                                                                                                                                                                                                                    organism="Homo sapiens"
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                                     2 GGACCCTAGAGGCAAA 17
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CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio:llnl.gov/bbrp/image/image.html
Insert Length: 574 Std Error: 0.00

Seq primer: -40UP from Gibco

High quality sequence stop: 409.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
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Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCT-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
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/mol type="mRNA"
/db_xref="taxon:9606"
/clome="TMAGE:2379150"
/lab_host="DH10B"
                                                                                                                                                                                                   /clone lib="NCI GGAP Kid11"
/note="Organ: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Eco RI; a modified polylinker; Site 1: Not I; Site 2: Eco RI; Plasmid DNA from the normalized library NCI GGAP Kid3 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 132336-1323911, 1456007-1456775, and
                                                                                                                                                                    Fatima Bonaldo.
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Pred. No.
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Pred. No.
                                                                          Mismatches
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sapiens cDNA clone IMAGE:2379150 3′,
                                                                                        DB 1;
2e+03;
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2e+03;
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RESULT 16
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                                                                        Homo sapiens (human)
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qg55a08.xl Soares_testis_NHT Homo sapiens cDNA clone

3' similar to gb:M58603 NUCLEAR FACTOR NF-KAPPA-B P10
                                                                                                                                                                                                                                                                                                                                                          2 GGACCCTAGAGGCAAAG 18
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Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T. Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett, G.L., Heaton, M.P., Laegreid, W.W., Rohrer, G.A., Chitko-McKown, C.G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J. and Keele, J.W.

Sequence 'evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle genome Res. 11 (4), 626-630 (2001)
                                                                                                                                    (HUMAN);, mRNA sequenc
AI208178
AI208178.1 GI:3770120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 402 762 4366
Fax: 402 762 4390
Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and alt_trimmed with phred
v0.980904.e. Vector identified by cross_match with the -minscore
Hominidae; Homo.
1 (bases 1 to 538)
                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                     TSE
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Seq primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FORWARD: AGGAAACAGCTATGACCAT
BACKWARD: GTTTTCCCAGTCACGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Smith TPL
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488200 MARC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Tvector: pCMV SPORT6; Site 1: NotI; Site 2: SalI; Library made from pooled tissue from testis, thymus, semitendonosus muscle, longissimus muscle, pancreas, adrenal, and endometrium."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="pooled"
/lab_host="DH10B"
/clone_lib="MARC 2BOV'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="mRNA"
/db_xref="taxon:9913"
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AG498153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
                                                                                                  Contribution of Asian mouse subspecies Mus musculus molossinus to genomic constitution of strain C57BL/6J, as defined by BAC-end sequence-SNP analysis
                                                                                                                                               Ezawa, K., Saitou, N., Hattori, M., Sakaki, Y., Shiroishi, T.
                                                                                                                                                                                                                                    AG498153
AG498153.1 GI:48205383
AG498153.1 GI:48205383
AG498153.1 GI:48205383
AG498153.1 GI:48205383
AG498153.1 GI:48205383
AG498153.1 GI:48205383

Mus musculus molossinus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             National Cancer Institute, Cancer Genome Anatomy F
Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
CDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                            AG498153
Mus musculus molossinus
Submitted (17-NOV-2003) Masahira Hattori,
                                                                                                                                                                                                                       Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Insert Length: 639 Std Error:
Seq primer: -40UP from Gibco
High quality sequence stop: 431.
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Insert Length: 639 Std Error: 0.00
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                              Hattori, M., Toyoda, A., Noguchi, H., Kojima, T.
                                                                                  Genome Res. 14 (12),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                   Direct Submission
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                                                                    .5574823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGACCCTAGAGGCAAA 17
                                              (bases 1 to 769)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="Soares_testis_NHT"
/notes_Tvector: pT7T3D-Pac (Pharmacia) with a modified
/notes_Tvector: pT7T3D-Pac (Pharmacia) with a modified
polylinker; Site_1: Not I; Site_2: Eco RI; lst strand cDNA
was prepared from mRNA obtained from Clontech
was prepared from mRNA obtained with a Not I - oligo(dT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:1839062"
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DNA,
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The Institute of Physical
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                                   and
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                                   Sakaki, Y
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genomic survey
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t5, and was
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BE503972
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Query Match
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1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa, 230-0045,
(B-mail:hattori@gsc.riken.jp, URL:http://hgp.gsc.riken.go.jp/,
Tel:81-45-503-911, Fax:81-45-503-9170)
Clones are derived from the mouse BAC library MSMg01. For BAC
Clones are derived from the mouse BAC library MSMg01. For BAC
library availability, please contact Kuniya Abe (abe@rtc.riken
Tsukuba Institude, Bio Resource Center,
The Institude of Physical and Chemical Research (RIKEN) 3-1-1
Koyadai, Tsukuba, 305-0074 Japan
phone: 81-298-36-9189, fax: 81-298-36-9199
e-mail: abe@rtc.riken.jp
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hz54a03.x1 NCI_CGAP_Lu24
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                                                                                                               Seq primer: -40UP from Gibco High quality sequence stop: 1
                                                                                                                                                         CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.B. Consortium/LLNL, send email to:

info@image.llnl.gov
                                                                                                                                                                                                                                                                                     Email: cgapbs-remail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                             Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA sequence.
BE503972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R.Site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R.Site 1 ':
                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases i to 123)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BE503972.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IBRARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      iomo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GACCCTAGAGGCAAAG 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GACCCTAGAGGCAAAG 18
                                                                                             quality sequence stop: 104.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue type="mixture of kidney and spleen"
/clone_lib="MSMg01 Mouse Male BAC Library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/sub_species="molossinus"
/db_xref="taxon:57486"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            clone="MSMg01-397021.T7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           . .769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GI:9706380
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EcoRI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bp mRN
o sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 10; 1
2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       molossinus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear EST 04-AUG-2000 clone IMAGE:3211756 3',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                        Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mg01. For BAC
(abe@rtc.riken.jp).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                        Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Query Match
Best Local Similarity
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                   source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   51 GGACCCTAGAGGCCAAG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                                                         www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 236 Std Error: 0.00
Seq primer: -40Up from Gibco.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
AI349272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AI349272
ta75c07.x2 NCI_CGAP_HSC2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .
G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tissue Procurement: Herbert Morse, M.D., Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 143)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AI349272.1 GI:4086478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA Library Preparation: David B. Krizman, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ph.D.
                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:2049900"
/tissue_type="stem_cell_34+/38+"
/dev_stage="adult"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clome_lib="NCI_CGAP_Lu24"
/clome_lib="NCI_CGAP_Lu24"
/note="Organ: lung; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Plasmid DNA from the normalized library NCI_CGAP_Lu5 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified CDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1414920-141799] and 1520904-1522439). Subtraction by Bento
                                                                                                 /clone lib="NCI_CGAP_HSC2"
/note="Organ: bone marrow; Vector: pAMP1; mRNA made from bone marrow, stem cells 34+/38+, cDNA made by oligo-dT priming. Directionally cloned. Size-selected on agarose gel, average insert size 400 bp. Primary library, non-amplified."
                                                                                             non-amplified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Soares and M. Fatima Bonaldo.
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/tissue_type="carcinoid"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   85.6%;
94.1%;
  85.6%;
94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
  Score 15.4; DB 1;
Pred. No. 3.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.4; DB 2;
Pred. No. 3.6e+03;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    143 bp mkNA
Homo sapiens cDNA
                                                                                                                   Size-selected on agarose Primary library,
                        Length 143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear EST 16-FEB-1999 clone IMAGE:2049900 3',
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Emmert-Buck,
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                      RESULT 21
AA234751 .
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JOURNAL COMMENT

FEATURES

ORIGIN

AA234751

177 bp

mRNA

linear

EST 03-MAR-1997

REFERENCE AUTHORS

TITLE

VERSION KEYWORDS

ACCESSION DEFINITION

SOURCE

ORGANISM

RESULT 19 AI349272/c

뭐 á ORIGIN

Matches

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JOURNAL COMMENT
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AUTHORS
TITLE
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KEYWORDS
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AA654607
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                                                                                    Matches
                                                                                                      Query Match
Best Local 9
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  65
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                                         2 GGACCCTAGAGGCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  M.D., Michael Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                High quality sequence stop: 142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  www-bio.llnl.gov/bbrp/image/image.html
Seq.primer: -40m13 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1997)
Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nt59e02.s1 NCI_CGAP_Pr3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA654607.1 GI:2590761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (bases 1 to 170)
                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                               histologically-determined to be fully malignant prostate cancer cells. Double-stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                            /clone_lib="NCI_CGAP_pr3"
/note="Vector: pAMPIO; Site_1: Not1; Site_2: EcoRI; 1st
strand cDNA was primed with oligo(dT)17 on 50 ng of
DNAse-treated, total cellular RNA obtained from
5,000-10,000 microdissected cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'dev_stage="45 years old"
'lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clone="IMAGE:1202810"
                                                                                                      85.6%;
94.1%;
81
                                                                                    <u>,</u>
                                                                                                      Score 15.4; DB 1;
Pred. No. 3.8e+03;
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                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNA
                                                                                                                           Length 170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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                                                                                                                                                                                                                                                                        RESULT 22
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                             53
                                                                                                                                                                AI099443

205 bp mRNA linear EST 20-AUG-1998
ue30f06.xl Sugano mouse liver mlia Mus musculus cDNA clone
IMAGE:1481891 3' similar to SW:ALBU_RAT P02770 SERUM ALBUMIN
                                                                                                                                                                                                                                                                                                                                                                                             N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The WashU-Merck EST Project Unpublished (1995) Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 177)
Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba;T., Le,M., Lennon,G., Marra,M.,
Parsons,J., Rifkin,L., Rohlfing;T., Soares,M., Tan,F.,
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            zs38c05.sl Soares_NhHMPu_S1 Homo sapiens cDNA clone IMAGE:687464
3', mRNA sequence.
AA234751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Seq primer: -41ml3 fwd. ET from Amersham High quality sequence stop: 157.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (human)
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                                                Mus musculus (house mouse)
                                                                                                   AI099443.1
                                                                                                                                                PRECURSOR. [1] ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                             GGACCCTAGAGGCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Soares_NhHMPu_Sl"
/note="Organ: mixed (see below); Vector: pT7T3D-Pac
/note="Organ: mixed (see below); Vector: pT7T3D-Pac
(Pharmacia) with a modified polylinker; Site_1: Not I;
Site_2: Eco RI; Equal amounts of plasmid DNA from three
normalized libraries (melanocyte 2NbHM, pregnant uterus
NbHPU, and fetal heart NbHH19W) were mixed, and ss circles
were made in vitro. Following HAP purification, this DNA
was used as tracer in a subtractive hybridization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  340488-345479, and 484488-489479."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="Pooled human melanocyte, fetal heart,
pregnant uterus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ocation/Qualifiers
                                                                                                   GI:3448968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .one="IMAGE:687464"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       _xref="GDB:5591614"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.4; DB 1;
Pred. No. 3.8e+03;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                     27
                                                                                                                                                                                                                     BI493189 215 bp ml df98e07.wl Morton Fetal Cochlea Homo IMAGE: 2540629 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T., Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CGGACCCTAGAGGCAAA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: custom primer used High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Marra M/Mouse EST Project
WashN-HHMI Mouse EST Project
WashIngton University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Lo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                        Homo sapiens
                                                                                                                                                                          BI493189.1 GI:15332533
                                                                                                                                                                                                      BI493189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The WashU-HHMI Mouse EST Project
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                                   Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                               iomo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                  CGGTCCCTAGAGGCAAA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 205)
       (bases 1 to 215)
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314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clome_lib="Sugano mouse liver mlia"
/clome_lib="Sugano mouse liver mlia"
/note="Organ: liver; Vector: pwB18S-Fi3; Site_1: DraIII
(CACTGTGTGT); Site_2: DraIII (CACCATGTG); 1st strand cDNA
was primed with an oligo(dT) primer
[ATGTGGCCTTTTTTTTTTTTTTTTTTT]; double-stranded cDNA was
ligated to a DraIII adaptor [TGTTGGCCTACTGG] cdlqested
and cloned into distrinct DraIII sites of the pwB18S-Fi3
vector (5' site CACTGTGTG, 3' site CACCATGTG). XhoI should
be used to isolate the cDNA insert. Size selection was
performed to exclude fragments <1.5kb. Library
constructed by Dr. Sumio Sugano (University of Tokyo
Institute of Medical Science). Custom primers for
sequencing: 5' end primer CTTCTGCTCTAAAAGCTGCG and 3' end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequencing: 5' end primer CTT primer CGACCTGCAGCTCGAGCACA."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /dev_stage="adult"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'clone="IMAGE:1481891"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             strain="C57BL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          85.6%;
94.1%;
                                                     Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.4; DB 1; Length 205; Pred. No. 3.8e+03; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           quality
                                                                                                                                                                                                                                                                                mRNA
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                                                                                                                                                                                                                                                      sapiens
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linear as cDNA

EST 28-AUG-2001 clone

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                          28
Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville,
                                                                 Zhao,S., Nierman,W., Malek,J., Shatsman,S., Al Tsegaye,G., Geer,K., Krol,M., Shvartsbeyn,A., Russell,D., de Jong,P. and Fraser,C.M. Mouse BAC End Sequences from Library RPCI-24 Unpublished (1999)
Other_GSSs: RPCI-24-236H6.TV
                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Euarchontoglires; Glires; Rodent Sciurognathi; Muroidea; Muridae; Murinae; Mus. 1 (bases 1 to 241)
                                                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                                                                                                      genomic survey sequence
BH084942
                                                                                                                                                                                                                                                                                                                                       RPCI-24-236H6.TJ RPCI-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and Morton,C.C.
Isolation of novel and known genes from a human fetal cochlear cDNA
library using subtractive hybridization and differential screening
                                                                                                                                                                                                                                                      Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                         BH084942.1
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Seq primer:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA sequencing and analyses were performed by National Institutes of Health Intramural Sequencing Center (NISC; see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reproductive Biology
Brigham and Women's Hospital
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMAGE Consortium (info@image.llnl.gov)
Plate: LLAM6328 row: I column: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Departments of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Morton, C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  http://www.nisc.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bmail: ccmorton@bics.bwh.harvard.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Robertson, N.G., Khetarpal, U., Gutierrez-Espeleta, G.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This clone is available royalty-free through LLNL; contact the MAGE Consortium (info@image.llnl.gov) for further information.
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617 738 6996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="cochlea"
/dev_stage="16-22 week fetus"
/lab_host="SOLR cells (kanamycin_resistant)"
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mol_type="mRNA"
                                                                                                                                                                                                                                                                                         GI:14904539
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Pred. No. 3.8e+03;
0; Mismatches 1;
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   MD 20850, USA
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                                                                                                                                                                                                                     Euteleostomi;
                                                                                                                                                                                                                                                                                                                                      RPCI-24-236H6,
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                                                                                                                                                   Levins, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16;
                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 258)
Robertson, N.G., Khetarpal, U., Gutierrez-Espeleta, G.A., Bieber, F.R. and Morton, C.C.
Isolation of novel and known genes from a human fetal cochlear cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   258 bp
df98e07.yl Morton Fetal Cochlea
IMAGB:2540629 5', mRNA sequence
BI493190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CGGACCCTAGAGGCAAA 17
                                                                                                                                                                                       75 Francis Street, Harvard Medical School, Tel: 617 732 7980
Fax: 617 738 6996
                                                                                                                                                                                                                                                                                                                                                           library using subtractive hybridization Genomics 23, 42-50 (1994) 7829101
Seq primer: M13RP1
                                          http://www.nisc.nih.gov).
This clone is available royalty-free through LLNL; contact the ThAGE Consortium (info@image.llnl.gov) for further information
                                                                                                                  DNA sequencing and analyses were performed of Health Intramural Sequencing Center (NIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (human)
                        Plate: LLAM6328
                                                                                                                                                                   Email: ccmorton@bics.bwh.harvard.edu
                                                                                                                                                                                                                                                               Brigham and Women's
                                                                                                                                                                                                                                                                                        Reproductive
                                                                                                                                                                                                                                                                                                         Departments of Pathology and Obstetrics,
                                                                                                                                                                                                                                                                                                                                     Contact: Morton, C. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BI493190.1 GI:15332534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seq primer: SP6
Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC (Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC page: http://ww .tigr.org/tdb/bac_ends/mouse/bac_end_intro.html plate: 236 row: H column: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: szhao@tigr.org
Clones are derived from the mouse BAC library RPCI-24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 301 838 0200 Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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/clone_lib="RPCI-24"
/clone_lib="RPCI-24"
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1 cloning vector at the library was cloned in the pTARBAC1 cloning vector at the BamH1 sites using MboI partially digested male C57BL/6J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Mus musculus"
/mol_type="qenomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'db_xref="taxon:10090"
'clone="RPCI-24-236H6"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Sutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                     Biology
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94.1%;
                        row: I column: 14
reverse primer
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Pred. No. 3.9e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                     BACKWARD: GTTTTCCCAGTCACGACG
Plate: 92 row: B column: 1
Seq primer: ATTTAGGTGACACTATAG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sus scrofa (pig)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and alt trimmed with phred
v0.980904.e. Vector identified by cross_match with the -minscore 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Smith TPL USDA, ARS, US Meat Animal Research Center PO Box 166, Clay Center, NE 68933-0166, US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fahrenkrug,S.C., Smith,T.P.L., Freking,B.A., Vallet,J., Wise,T., Rohrer,G.A., Pertea,G., Quackenbush,J. and Keele,J.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 402 762 4366
Fax: 402 762 4390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CR PRimers
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/tissue type="cochlea"
/dev stage="16-22 week fetus"
/lab_host="SOLR cells (kanamycin resistant)"
/clone_lib="Morton Fetal Cochlea"
/clone_lib="Morton Fetal Cochlea"
/note="Organ: ear; Vector: pBluescript SK-; Site_1: EcoRI;
/note="Organ: ear; Vector: pBluescript SK-; Site_1: EcoRI;
/note="Organ: ear; Vector: pBluescript SK-; Site_1: EcoRI;
/note="Cryan: ear; Vector: pBluescript SK-; Site_1: EcoRI;
/note="Organ: ear; Vector: pBluescript SK-; Site_1: EcoRI;
/note="Organ: ear; Vector: Organ: PBluescript SK-; Site_1: EcoRI;
/stage="Cochlea"
/file="Cochlea"
/file="
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome 13 (8), 475-478 (2002)
                                                                                                                                                                /organism="Sus scrofa"
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/db_xref="taxon:9823"
/tissue_type="pooled"
/lab_host="DH10B"
                                                            /clone_lib="MARC 1PIG"
/note="Vector: pCMV SPORT6; Site_1: NotI; Site_2:
Library made from pooled tissue from day 11, 13, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                            location/Qualifiers
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94.1%;
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RESULT 28
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                                                           mRNA sequence
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nx62h05.si NCI_CGAP_Alvi
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seq primer: -4onl3 fwd. ET from Amersham.
Location/Qualifiers
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Unpublished (1997)
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                               AA747067.1 GI:2787025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                                                   Similarity
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
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sapiens (human)
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/note="Vector: pAMP10; mRNA made from alveolar
/note="Vector: pAMP10; mRNA made from alveolar
/nabdomyosarcoma, cDNA made by oligo-dT priming.
Non-directionally cloned. Size-selected on agarose gel,
average insert size 600 bp. Reference: Krizman et al.
(1996) Cancer Research 56:5380-5383."
                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:1266873"
/tissue_type="alveolar rhabdomyosarcoma"
/lab_host="DH10B"
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/mol_type="mRNA"
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Pred. No. 3.9e+03;
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sapiens cDNA clone IMAGE:1267125,
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sapiens cDNA clone IMAGE:1266873,
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AW087244/c
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Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                   Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michae
R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life
Technologies, Inc. cDNA Library Arrayed by: Christa Prange, The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                      NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                 Hominidae; Homo.
1 (bases 1 to 313)
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 371 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham.
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
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Mammalia; Eutheria;
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                                                         .M.A.G.E. Consortium DNA Sequencing by: Washington University
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DNA Sequencing by: Washington University Genome Sequencing
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/note="Vector: pAMP10; mRNA made from alveolar
rhabdomyosarcoma, cDNA made by oligo-dT priming.
Non-directionally cloned. Size-selected on agarose caverage insert size 600 bp. Reference: Krizman et
(1996) Cancer Research 56:5380-5383."
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/mol_type="mRNA"
/db_xref="taxon:9606"
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/tissue_type="alveolar rhabdomyosarcoma"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AUTHORS
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BM743266
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Kim, N.S., Haum, A., Sohn, B. Cheong, J.E., Sohn, B. Cheong, J.E., Sohn, B. Cheong, Y.S. 21C Frontier Korean EST Project 2001
                                                                                                                                                                                                                                                                                                                                                                                         Genome Research Center
Korea Research Institute of Bioscience & Biotechnology
52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                 Email: yongsung@mail.kribb.re.kr
Plate: 10 row: E column: 01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 K-EST0016364 S4SNU1 Homo sapiens cDNA clone S4SNU1-10-E01 5', mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGACCCTAGAGGCAAAG 18
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                                                                                                                                                                                                                                                                                                                             quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              quality sequence stop: 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
/note="Organ: Stomach; Vector: pTZ18RP1; Site 1: EcoR1; Site 2: NotI; The poly (A)+ RNA was dephosphorylated wibacterial alkaline phosphatase (BAP) and then decapped
                                                                                         /lab_
                                                                                                                                                                                       /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="S4SNU1-10-E01"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="NCI_CGAP_CO21";
/note="Organ; colon; Vector: pCMV-SPORT6; Site_1: Sal
Site_2: NotI; Cloned unidirectionally. Primer: Oligo
Normalized to Cot >500. Average insert size 1.04kb.
Normalized version of NCI_CGAP_Co18. Library constr
by Life Technologies."
                                                                  /clone_
                                                                                                           cell_type="Lympio"
                                                                                                                                                   tissue_type="Stomach"
                                                                                                                                                                              "Bex="M"
                                                                                                                                                                                                                                        organism="Homo sapiens"
|mol_type="mRNA"
                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            clone="IMAGE:2582708"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tissue_type="moderately differentiated adenocarcinoma"
lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GI:19064595
                                                                                         host="Top10F'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  85.6%;
94.1%;
                                                                                                                               type="Lymphoblast-like"
                                                                  lib="S4SNU1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.,
E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                254
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                             with
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Query Match 85.6%; Score 15.4; DB 3; Length 326; Best Local Similarity 94.1%; Pred. No. 4e+03; Matches 16; Conservative 0; Mismatches 1; Indels with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli T0plUF by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."

. 0;

Gaps

0;

ORIGIN

Search completed: April 15, 2006, 18:51:14 Job time: 3827 secs